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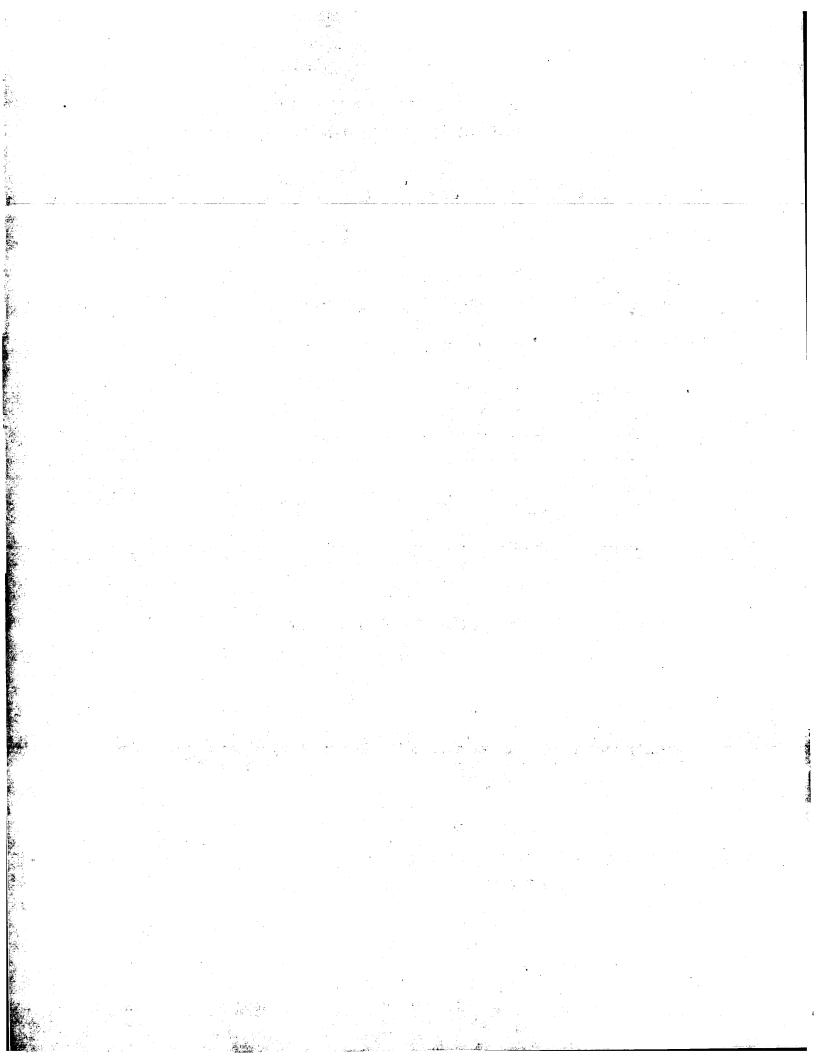
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PATENT COOPERATION TREATY

From the INTERNATIONAL BUREAU

PCT

NOTIFICATION OF ELECTION

(PCT Rule 61.2)

To:

United States Patent and Trademark Office (Box PCT) Crystal Plaza 2 Washington, DC 20231 ETATS-UNIS D'AMERIQUE

Date of mailing (day/month/year) 25 February 1998 (25.02.98)	in its capacity as elected Office
International application No. PCT/NL97/00345	Applicant's or agent's file reference PCT 0589
International filing date (day/month/year) 19 June 1997 (19.06.97)	Priority date (day/month/year) 20 June 1996 (20.06.96)
Applicant VAN LEENGOED, Leonardus, Andrianu	s, Maria, Govardus et al

1.	The designated Office is hereby notified of its election made:
	X in the demand filed with the International Preliminary Examining Authority on:
	20 January 1998 (20.01.98)
	in a notice effecting later election filed with the International Bureau on:
2.	The election X was
	was not
:	made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).
-	

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland Authorized officer

R. Raissi

Telephone No.: (41-22) 338.83.38

Facsimile No.: (41-22) 740.14.35



INTERNATIONAL SEARCH REPORT

(PCT Article 18 and Rul s 43 and 44)

Applicant's or agent's file reference	FOR FURTHER see Notification of	Transmittal of International Search Report 20) as well as, where applicable, item 5 below.
PCT 0589	ACTION	
International application No.	International filing date (day/month/year)	(Earliest) Priority Date (day/month/year)
PCT/NL 97/00345	19/06/1997	20/06/1996
Applicant		
KOSTER. Henk W. et al.		
KOSTER, Henk W. et al.		
This International Search Report has bee according to Article 18. A copy is being tr	on prepared by this International Searching Auth ansmitted to the International Bureau.	ority and is transmitted to the applicant
This International Search Report consists X It is also accompanied by a cop	s of a total of sheets. by of each prior art document cited in this report.	·
1. X Certain claims were found ur	nsearchable (see Box I).	
2. Unity of invention is tacking (see Box II).	
3. X The international application of international search was carried	ontains disclosure of a nucleotide and/or amin d out on the basis of the sequence listing	o acid sequence listing and the
	d with the international application.	
· —	nished by the applicant separately from the inte	
	but not accompanied by a statement to the matter going beyond the disclosure in the	e effect that it did not include international application as filed.
Tra	anscribed by this Authority	
4. With regard to the title, the	e text is approved as submitted by the applicant.	
T the	e text has been established by this Authority to r	ead as follows:
IL-6 AND IL-6-RECEPTO ACTIVITY	OR DERIVED PEPTIDES HAVING II	6 ANTAGONISTIC OR AGONISTIC
5. With regard to the abstract,		
	e text is approved as submitted by the applicant	
1 B	e text has been established, according to Rule 3 ox III. The applicant may, within one month from earch Report, submit comments to this Authority	the date of mailing of this international
6. The figure of the drawings to be pu	blished with the abstract is:	
Figure No as	suggested by the applicant.	None of the figures.
	ecause the applicant failed to suggest a figure.	
be	ecause this figure better characterizes the inven	ion.

International application No. PCT/NL 97/00345

INTERNATIONAL SEARCH REPORT

Box I Obs rvati ns where c rtain claims were found uns archable (Continuation of it in 1 of first she t)	
This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:	
1. X Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:	
see FURTHER INFORMATION sheet PCT/ISA/210	
2. Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:	
Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).	
B x II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)	
This International Searching Authority found multiple inventions in this international application, as follows:	
As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.	
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.	
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:	
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:	
Remark on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.	

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Remark: Although claim(s) 18 and 19 (both partially, as far as an in vivo method is concerned) are directed to a diagnostic method practised on the human/animal body, and although claims 15 and 22 (both completely) and claim 16 (partially, as far as an in vivo method is concerned) are directed to a method of treatment of the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.





According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 CO7K A61K GO1N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO	BE RELEVANT

Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
х	DATABASE WPI Week 9607	1,3,6, 14-16
	Derwent Publications Ltd., London, GB; AN 96-065477	
	XP002027134 & JP 07 324 098 A (DAICEL CHEM. IND. LTD. ET AL.) , 12 December 1995	
	see abstract	
X	DATABASE WPI Week 9607	1-6, 14-16
	Derwent Publications Ltd., London, GB; AN 96-065476	
	XP002027135 & JP 07 324 097 A (DAICEL CHEM. IND. LTD. ET AL.) , 12 December 1995	
	see abstract	
	-/	

X Further documents are listed in the continuation of box C.	Patent family members are listed in annex.		
Special categories of cited documents : A* document defining the general state of the art which is not considered to be of particular relevance.	T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention		
"E" earlier document but published on or after the international filing date "L" document which may throw doubts on priority claim(s) or	"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone		
which is cited to establish the publication date of another citation or other special reason (as specified)	"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such docu-		
O document referring to an oral disclosure, use, exhibition or other means *P* document published prior to the international filing date but later than the priority date claimed	document is combined with one of more other such docu- ments, such combination being obvious to a person skilled in the art. "&" document member of the same patent family		
Date of the actual completion of the international search	Date of mailing of the international search report		
20 October 1997	.1 0. 11. 97		
Name and mailing address of the ISA	Authorized officer		
European Patent Office, P.B. 5818 Patentiaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016	Nooij, F		

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Category °	ation) DOCUMENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
x	DATABASE WPI Week 9035 Derwent Publications Ltd., London, GB; AN 90-266224 XP002027136 & JP 02 188 600 A (CHUGAI PHARMACEUTICAL KK), 24 July 1990 see abstract	1,3,6, 14-18
X	WO 95 04075 A (MEDVET SCIENCE PTY. LTD.) 9 February 1995 see examples 15,16 see seq. id. nos 17,36,37 see table 2	1-7, 14-16,21
(EP 0 426 857 A (KURARAY CO. LTD.) 15 May 1991 see claim 7	2,3,6,7, 16
X	US 5 210 075 A (SCHOLZ ET AL.) 11 May 1993 see examples	1-5,12, 14-16
x	C. MORTON ET AL.: "Solution structure of synthetic peptides corresponding to the C-terminal helix of interleukin-6." EUROPEAN JOURNAL OF BIOCHEMISTRY, vol. 219, no. 1-2, 15 January 1994, BERLIN, GERMANY, pages 94-107, XP000645297 see page 98, left-hand column, line 50 - line 61 see page 105, right-hand column, line 20 - page 106, left-hand column, line 4	1,3,4,6,
Α	M. KALAI ET AL.: "Participation of two Ser-Ser-Phe-Tyr repeats in interleukin-6 (IL-6) binding sites of the human IL-6 receptor." EUROPEAN JOURNAL OF BIOCHEMISTRY, vol. 238, no. 3, June 1996, BERLIN, GERMANY, pages 714-723, XP000645294 see table 1	12

ERNATIONAL SEARCH REPORT

Information on patent family members

Internationa	Application No	
PCT/NL	97/00345	

Patent document cited in search report			Publication date	
WO 9504075 A	09-02-95	AU 7341494 A CA 2168261 A EP 0715633 A JP 9501154 T	28-02-95 09-02-95 12-06-96 04-02-97	
EP 426857 A	15-05-91	CA 2026881 A WO 9009396 A US 5171837 A	09-08-90 23-08-90 15-12-92	
US 5210075 A	11-05-93	NONE		

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INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification 6:

C07K 14/54, 14/715, 16/24, 16/28, 16/42, A61K 38/17, 38/20, 39/395, G01N 33/68, 33/577

A1 (

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24 December 1997 (24.12.97)

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19 June 1997 (19.06.97)

(30) Priority Data;

96201720.8 20 June 1996 (20.06.96) (34) Countries for which the regional or

.96) EP

AT et al.

(71) Applicant (for all designated States except US): KOSTER, Henk, Wilhelmus [NL/MC]; 1607 Parc Saint Roman, MC-98000 Monte Carlo (MC).

international application was filed:

(72) Inventors; and

- (75) Inventors/Applicants (for US only): VAN LEENGOED. Leonardus, Andrianus, Maria, Govardus [NL/NL]; Oostrandpark 6, NL-8212 AN Lelystad (NL). HOEBE, Kasper, Hubertus, Nicolaas [NL/NL]; Tolsteegplantsoen 46, NL-3523 AN Utrecht (NL). HOEBE, Kasper, Hubertus, Nicolaas [NL/NL]; Karveel 10-04, NL-8231 AP Lelystad (NL).
- (74) Agent: SMULDERS, Th., A., H., J.; Vereenigde Octrooibureaux, Nicuwe Parklaan 97, NL-2587 BN The Hague (NL).

(81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LY, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, ARIPO patent (GH, KE, LS, MW, SD, SZ, UG, ZW), Burasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DB, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CP, CG, CI, CM, GA, GN,

Published

With international search report.

ML, MR, NE, SN, TD, TG).

Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.

(54) Title: IL-6 AND IL-6-RECEPTOR DERIVED PEPTIDES HAVING IL-6 ANTAGONISTIC OR AGONISTIC ACTIVITY

(57) Abstract

The invention relates to IL-6 and IL-6-receptor derived peptides having IL-6 agonistic or antagonistic activity. The peptides are at least 5 amino acids long and are selected from one of the following amino acid sequences: RYILDGISALRK, STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQMQLSCFRKSPLSNVVC, PRSTPSLTTKAVLLVRKFQNS, MCVASSVGSKFSK-TQTFQGC, PEKPKNLSCIVNEGKKMRCBWDGGR, NFTLKSEWATHKFADCKAKRDTPTS, WVEABNALGKVTSDH, EWGPRSTP-SLTTKAVLLVRKFQNSPAED or PVYKVKPNPPHNLSVIN. Selected peptides and combinations of selected peptides can be used in the treatment, prevention, detection, or diagnosis of IL-6 related disease and can be used to clear blood or bloodproducts of IL-6 or IL-6 receptor molecules.

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VEREENIGDE DEN HAAGNO 2.... NO. 6357 P. 5 Continuation of Box No. III WURTHER APPLICANTS AND/OR (FURTHER) INVENTORS If none of the following sub-boxes is used, this sheet is not to be included in the request. Name and address: (Family name followed by given name: for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (i.e. country) of residence if no State of residence is indicated below.) This person is: applicant only Kasper, Hubertus Nicolaas Hoebe Tolsteegplantsoen 46 applicant and inventor 3523 AN Utrecht the Netherlands inventor only (If this check-box is marked, do not fill in below.) State (i.e. country) of nationality: State (i.e. country) of residence: NL NL This person is applicant all designated all designated States except the United States of America the United States of America only the States indicated in for the purposes of: States the Supplemental Box Name and address: (Family name followed by given name: for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (i.e. country) of residence if no State of residence is indicated below.) This person is: applicant only Meloen, Robert Hans Karveel 10-04 applicant and inventor 8231 AP Lelystad the Netherlands inventor only (If this check-box is marked do not fill in below.) State (i.e. country) of nationality: State (i.e. country) of residence: NL NL the United States of America only the States indicated in the Supplemental Box This person is applicant all designated States all designated States except the United States of America for the purposes of: Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (i.e. country) of residence if no State of residence is indicated below.) This person is: applicant only applicant and inventor inventor only (If this check-box is marked, do not fill in below.) State (i.e. country) of nationality: State (i.e. country) of residence: This person is applicant the States indicated in the Supplemental Box all designated all designated States except the United States of America the United States of America only for the purposes of: Name and address: (Family name followed by given name; for a legal entity, full official designation. The address must include postal code and name of country. The country of the address indicated in this Box is the applicant's State (i.e. country) of residence if no State of residence is indicated below.) This person is: applicant only applicant and inventor inventor only (If this check-box is marked, do not fill in below.) State (i.e. country) of nationality: State (i.e. country) of residence: This person is applicant for the purposes of: al designated all designated States except the United States of America the United States

of America only

the States indicated in

the Supplemental Box

P 6

Box No.Y DESIGN

DESIGNATION STATES

The following designations are hereby made under Rule 4.9(a) (mark the applicable check-boxes: at least one must be marked):
Regional Patent

X	AP	ARIPO Patent: KE Kenya, LS Lesotho, MW Malawi, SD Sudan, SZ Swaziland, UG Uganda, and any other State which is a Contracting State of the Harare Protocol and of the PCT
		is a Contracting State of the Harare Protocol and of the PC I

- EA Eurasian Patent: AM Armenia, AZ Azerbaijan, BY Belarus, KG Kyrgyzstan, KZ Kazakstan, MD Republic of Mold va, RU Russian Federati n, TJ Tajikistan, TM Turkmenistan, and any other State which is a Contracting State of the Eurasian Patent C nvention and of the PCT
- EP European Patent: AT Austria, BE Belgium, CH and LI Switzerland and Liechtenstein, DE Germany, DK Denmark, ES Spain, FI Finland, FR France, GB United Kingdom, GR Greece, IE Ireland. IT Italy, LU Luxembourg, MC Monaco. NL Netherlands, PT Portugal, SE Sweden, and any other State which is a Contracting State of the European Patent Convention and of the PCT
- OAPI Patent: BF Burkina Faso, BJ Benin, CF Central African Republic, CG Congo, CI Côte d'Ivoire, CM Cameroon, GA Gabon, GN Guinea, ML Mali, MR Maurimia, NE Niger, SN Senegal, TD Chad, TG Togo, and any other State which is a member State of OAPI and a Contracting State of the PCT (if other kind of protection or treatment desired, specify on dotted line)

Nation	tal P	ateut (if other kind of protection or preaument desired.	speci	ify on	dotted line):
X	ΑL	Albania	X	LU	Luxembourg
X	AM	Armenia	X	LY	Larvia
X	ΑT	Austria	X	MD	Republic of Moldova
X	ΑU	Australia	X	MG	Madagascar
X	AZ.	Azerbaijan	\mathbb{X}	МK	The former Yugoslav Republic of Macedonia
X	BA	Bosnia and Herzegovina			
X	BB	Barbados	X	MN	Mongolia
X	BG	Bulgaria	X	MW	Malawi
$\overline{\mathbf{x}}$	BR	Brazil :	X	MX	Mexico
$\overline{\mathbf{x}}$	BY	Belarus	X	OK.	Norway
X	CA	Canada	$\overline{\mathbf{x}}$	NZ	New Zealand
X	CH:	and LI Switzerland and Liechtenstein	$\overline{\mathbf{x}}$	PL	Poland
X	CN	China	\mathbf{x}	PT	Portugal
\mathbf{x}	CU	Cuba	$\overline{\mathbf{x}}$	RO	Romania
X	CZ	Czech Republic	X	RU	Russian Federation
X	DE	Germany	X	SD	Sudan
X	DK	Denmark		SE	Sweden
X	EE	Estonia	$\overline{\mathbf{x}}$	SG	Singapore
X	ES	Spain	$\overline{\mathbf{x}}$	SI	Slovenia
\boxtimes	FI	Finland		SK	Slovakia
図	GB	United Kingdom	$\overline{\mathbb{Z}}$	ŢJ	Tajikistan
X	GE	Georgia	区	TM	Turkmenistan
X	ИŲ	Hungary	$\overline{\mathbb{K}}$	TR	Turkey
X	IL	Israel		TT	Trinidad and Tobago
X	IS	Iceland	X	UA	Ukraine
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X	KP	Democratic People's Republic of Korea	X	UZ	Uzbekistan
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X	KR	Republic of Korea	~		
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X	LC	Saint Lucia	issue		f this sheet:
図	_	Sri Lanka			Ghana
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	LS	Lesotho	_		Zimbabwe
E	LT	Lithuania	<u> </u>	بدج	Sierra Leone

In addition to the designations made above, the applicant also makes under Rule 4.9(b) all designations which would be permitted under the PCT except the designation(s) of

The applicant declares that those additional designations are subject to confirmation and that any designation which is not confirmed before the expiration of 15 months from the priority date is to be regarded as withdrawn by the applicant at the expiration of that time limit. (Confirmation of a designation consume of the filling of a nounce specifying that designation and the payment of the designation and confirmation fees. Confirmation must reach the receiving Office within the 15-month time (limit.)

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The priority of the following es	ulicr application(s) is hereby claim	urther priority claims are indicated in	the Supplemental E
I Country	appreacion(s) is necesy claim	ed:	
(in which, or for which, the application was filed)	Filing Date (day/monsh/year)	Application No.	Office of f (only for regi international a
item (1) EP	20 June 1996	96201720.8	inconstitutional ap
item (2)	20. 06. 1996	30201720.8	
item (3)			
The receiving Office is her Bureau a certified copy of	reby requested to prepare and trans the earlier application(s) identified	to be usued by the Office which for the purp mit to the International above as item(s):	oses of the present in
Box No. VII INTERNATION	AL SEARCHING AUTHORITY	Y	
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BUT OF requested and the Authorisis	ch (international, international-type or tow requested to base the international, ence to the relevant application (or the Date (day/month/year):	other) by the International Searching Auth search, to the extent possible, on the results translation thereof) or by reference to the Number:	ority has already bee of that earlier search search request:
EP	7 March 1997	9620172	8.01
Box No. VIII CHECK LIST			 -
This international application	annin Tining		
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International Searching Authority specified by the applicant:	ISA / 6.	Transmittal of search copy delayed until search fee is paid	\dashv

INTERNATIONAL SEARCH REPORT

International Ap PCT/NL 97/00345

A. CLASSIFICATION OF SUBJECT MATTER IPC 6 C07K14/54 C07K14/715

A61K38/17

A61K38/20

· C07K16/24 A61K39/395 C07K16/28 G01N33/68 C07K16/42 G01N33/577

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

CO7K A61K GO1N IPC 6

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the International search (name of data base and, where practical, search terms used)

C. DOCO	ALM IS CONSIDERED TO BE RELEVANT	

Oalegory *	Citation of document, with Indication, where appropriate, of the relevant passages	Relevant to claim No.
X	DATABASE WPI 🖒 / Week 9607	1,3,6, 14-16
	Derwent Publications Ltd., London, GB; AN 96-065477 XP002027134	
	& JP 07 324 098 A (DAICEL CHEM. IND. LTD. ET AL.) , 12 December 1995	
	see abstract	·
X	DATABASE WPI Week 9607	1-6, 14-16
,	Derwent Publications Ltd., London, GB; AN 96-065476	
	XP002027135 & JP 07 324 097 A (DAICEL CHEM. IND. LTD. ET AL.) , 12 December 1995	
	see abstract	

X Further documents are listed in the continuation of box C.

X

Patent family members are listed in annex.

- Special calegories of cited documents:
- *A* document defining the general state of the art which is not considered to be of perticular relevance
- earlier document but published on or after the international films date
- "L" document which may throw doubts an priority claim(s) or which is ofted to establish the publication date of another citation or other special reason (as specified)
- "O" document referring to an oral disolosure, use, exhibition or other means
- document published prior to the international filing date but later than the priority date claimed
- "I" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle of theory underlying the invention
- "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is laken alone
- document of particular relevance; the claimed invention cannot be considered to involve an inventive stap when the document is combined with one or more other such documents, such combination being covious to a person skilled
- "&" document member of the same patent family

Date of the actual completion of the international search

Oute of mailing of the international search report

20 October 1997

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Name and mailing address of the ISA

1

European Palent Office, P.B. 5818 Patentiann 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni, Fax: (+31-70) 340-3016

Authorized officer

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Form PCT/ISA/210 (execond sheet) (July 1992)

INTERNATIONAL SEARCH REPORT

PCT/NL 97/00345

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the relevant passages Relevant to claim No.
1,3,6, 14-18 MARMACEUTICAL
CE PTY. LTD.) 9
LTD.) 15 May 2,3,6,7,
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Internatio pplication No. PC1/NL 97/00345

INTERNATIONAL SEARCH REPORT

Boxi	Observations where c rtain claims were found uns archable (Continuation of item 1 of first sheet)
This Inte	mational Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. X	Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely: see FURTHER INFORMATION sheet PCT/ISA/210
· 2.	Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
з, 🔲	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II	Observations where unity of invention is tacking (Continuation of Item 2 of first sheet)
This Int	emational Searching Authority found multiple inventions in this international application, as follows:
:	
1.	As all required additional search less were timely paid by the applicant, this international Search Report covers all searchable claims.
: : 2	As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
э	As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4.	No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
fiern a i	The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.

NO. 6357

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International Ar Alton No PCT/NL 97/00345

INTERNATIONAL SEARCH REPORT

information on patent family members

Patent document cited in search report	Publication date	Paterd family member(s)	Publication date
NO 9504075 A	09-02-95	AU 7341494 A CA 2168261 A EP 0715633 A JP 9501154 T	28-02-95 09-02-95 12-06-96 04-02-97
EP 426857 A	15-05-91	CA 2026881 A WO 9009396 A US 5171837 A	09-08-90 23-08-90 15-12-92
US 5210075 A	11-05-93	NONE	•

From the INTERNATIONAL BUREAU PCT **NOTIFICATION OF THE RECORDING** OF A CHANGE SMULDERS, Th., A., H., J., c.s. Vereenigde Octrooibureaux (PCT Rule 92bis.1 and Nieuwe Perklaan 97 NL-2587 BN The Hague Administrative Instructions, Section 422) **PAYS-BAS** Date of mailing (day/month/year) 10 November 1997 (10.11.97) Applicant's or agent's file reference IMPORTANT NOTIFICATION **PCT 0589** International application No. International filing date (day/month/year) PCT/NL97/00345 19 June 1997 (19.06.97) 1. The following indications appeared on record concerning: X the applicant the inventor the agent the common representative State of Nationality State of Residence Name and Address NL MC KOSTER, Henk, W. Telephone No. Facsimile No. Teleprinter No. 2. The International Bureau hereby notifies the applicant that the following change has been recorded concerning: the person the name the address the residence the nationality State of Nationality State of Residence Name and Address KOSTER, Henk, Willielmus MC NL Telephone No. Facsimile No. Teleprinter No. 3. Further observations, if necessary: 4. A copy of this notification has been sent to: the receiving Office the designated Offices concerned the International Searching Authority the elected Offices concerned the International Preliminary Examining Authority other:

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland Authorized officer

Telephone No.: (41-22) 338.83.38

Martine Lee



Facsimile No.: (41-22) 740.14.35

From the INTERNATIONAL BUREAU					
PCT	To:				
NOTIFICATION OF THE RECORDING OF A CHANGE (PCT Rule 92bis.1 and Administrative Instructions, Section 422) Date of mailing (day/month/year)	SMULDERS, Th., A., H., J., c.s. Vereenigde Octro ibureaux Nieuwe Parklaan 97 NL-2587 BN The Hague PAYS-BAS				
01 October 1997 (01.10.97)					
Applicant's or agent's file reference PCT 0589	IMPORTANT NOTIFICATION				
International application No. PCT/NL97/00345	International filing date (day/month/year) 19 June 1997 (19.06.97)				
The following indications appeared on record concerning: X the applicant X the inventor	the agent the common representative				
Name and Address KASPER, Hubertus, Nicolaas, Hoebe	State of Nationality State of Residence NL NL Telephone No. Facsimile No. Teleprinter No.				
2. The International Bureau hereby notifies the applicant that to the person X the name the ad-	l I I I sho rogidasas I				
Name and Address HOEBE, Kasper, Hubertus, Nicolaas	NL NL Telephone No.				
	Facsimile No.				
	Teleprinter No.				
3. Further observations, if necessary:					
A copy of this notification has been sent to: X the receiving Office X the International Searching Authority the International Preliminary Examining Authority	the designated Offices concerned the elected Offices concerned other:				
The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland	Authorized officer Martine Lee				
Facsimile No.: (41-22) 740.14.35	Telephone No.: (41-22) 338.83.36				

pus. (uf ONTVANGEN From the INTERNATIONAL PRELIMINARY EXAMINING AUTHORITY PC70KT 1998 To: AMERSFOORT Koris SMULDERS, Th.A.H .h/naar TERMIJN NOTIFICATION OF TRANSMITTAL OF VEREENIGDE OCTROOIBUREAUX. THE INTERNATIONAL PRELIMINARY 1 3 JAT. 1998 Nieuwe Parklaan 97 EXAMINATION REPORT 2587 BN Den Haag Besniwsoni Buricht gezonder PAYS-BAS (PCT Rule 71.1) veoil. ลอก def. dd. Date of mailing 0 8, 10. 98 0589 (day/month/year) Applicant's or agent's file reference IMPORTANT NOTIFICATION PCT 0589 Priority date (day/month/year) International filing date (day/month/year) International application No. 20/06/1996 19/06/1997 PCT/NL97/00345 Applicant

- 1. The applicant is hereby notified that this International Preliminary Examining Authority transmits herewith the international preliminary examination report and its annexes, if any, established on the international application.
- 2. A copy of the report and its annexes, if any, is being transmitted to the International Bureau for communication to all the elected Offices.
- 3. Where required by any of the elected Offices, the International Bureau will prepare an English translation of the report (but not of any annexes) and will transmit such translation to those Offices.

4. REMINDER

KOSTER, Henk Wilhelmus et al.

The applicant must enter the national phase before each elected Office by performing certain acts (filing translations and paying national fees) within 30 months from the priority date (or later in some Offices) (Article 39(1)) (see also the reminder sent by the International Bureau with Form PCT/IB/301).

Where a translation of the international application must be furnished to an elected Office, that translation must contain a translation of any annexes to the international preliminary examination report. It is the applicant's responsibility to prepare and furnish such translation directly to each elected Office concerned.

For further details on the applicable time limits and requirements of the elected Offices, see Volume II of the PCT Applicant's Guide.

Name and mailing address of the IPEA/

European Palent Office D-80298 Munich

Tel. (+49-89) 2399-0 Tx 523656 epmu d

Fax: (-49-89) 2399-4465

Authorized officer

Peralt Lappas, R

Tel. (+49-89) 2399-8052



PATENT COOPERATION TREATY

PCT

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's or agent's file reference PCT 0589			FOR FURTHER AC	TION See	See Notification of Transmittal of International Preliminary Examination Report (PCT/IPEA/416)		
	plication	on No.	International filing date (day/m	onth/year)	Priority date (day/month/year)	,	
Wite in a series of the series			19/06/1997		20/06/1996		
			ational classification and IPC				
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Applicant							
	lenk \	Wilhelmus et al.					
						mining Authority	
1. This integral and is tr	rnatio ansmi	nal preliminary exa itted to the applican	mination report has been pre t according to Article 36.	pared by this in	ternational Preliminary Exa	initing Additions	
1			of 7 shéets, including this c				
⊠ Th wh be	is repo ich ha fore th	ort is also accompa we been amended is Authorlty (see R	nled by ANNEXES, i.e., she and are the basis for this rep ule 70.16 and Section 607 of	ets of the descrip ont and/or sheet the Administrati	otion, claims and/or drawing is containing rectifications n ive Instructions under the P	nade CCT).	
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3. This re	port co	ontains indications i	relating to the following items	:			
	Ø	Basis of the repor	t				
i n		Priority				sility	
III		Non-establishmer	nt of opinion with regard to no	ovelty, inventive	step and industrial applicat	July .	
IV		Lack of unity of in	vention			Landiaphitity	
V	☒	Reasoned statem citations and expl	ent under Article 35(2) with r anations supporting such sta	regard to novelty atement	, inventive step or industria	паррисаопку,	
VI		Certain documen	ts cited				
- VII		Certain defects in the international application					
VIII	×	Certain observati	ons on the international appl	ication			
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ı	Fa	_{IX.} (±49-89) 2399-446!	5	1			

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/NL97/00345

I.	Basi	s of th	report		•				
1.	This report has been drawn on the basis of (substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments.):								
	Description, pages:								
	1-16			as originally file	d				
	Clair	ms, No.	:						
	1-24		as received on		18/06/1998	with letter of	18/06/1998		
	Drav	wings, s	sheets:						
	1-5			as originally file	ed				
2	. The	amendı	ments hav	e resulted in the	cancellation of	of:			
		the des	cription,	pages:	,				
	×	the clai	ims,	Nos.:	1-23				
		the dra	wings,	sheets:					
3	3. This report has been established as if (some of) the amendments had not been made, since they have been considered to go beyond the disclosure as filed (Rule 70.2(c)):							n made, since they have been	
4	I. Add	ditional d	observatio	ns, if necessary:					

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INTERNATIONAL PRELIMINARY **EXAMINATION REPORT**

International application No. PCT/NL97/00345

V. R asoned statement under Article 35(2) with regard to novelty, inventive st p or industrial applicability; citations and explanations supporting such statement

1. Statement

Novelty (N)

Yes:

Claims CLAIMS 2, 6-12

No: '

Claims CLAIMS 1, 3-5, 13-24

Inventive step (IS)

Yes:

Claims CLAIMS 2, 6-11

No:

Claims CLAIM 12

Industrial applicability (IA)

Yes:

Claims CLAIMS 1-24 Claims

No:

2. Citations and explanations

see separate sheet

VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:

see separate sheet

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- 1.1 The documents cited in the ISR ar labell d D1 to D8.
- 1.2 The claims are allowable under Art.34(1)(b) PCT.

AD SECTION VIII:

- 2. Art. 6 EPC (clarity)
- 2.1 The words " long " in claims 6, 7 and 10 render the claims unclear, for obvious reasons.
- 2.2 The term in claim 20 " at concentrations relatively equivalent to bio assay " lacks clarity (Art. 6 PCT). The clear limits of this let us say range feature are not stated in the claim and different views of the limits of the range are conceivable for different skilled persons. The claim is subject to subjective interpretation and it therefore lacks clarity.

AD SECTION V.2.:

- 3. Art. 33(2) and (3) over D5
- 3.1 The peptides of the general formula of claim 1 of **D5**, when A is a peptide fragment of the formulas of claims 2 to 13 (the formulas in claims 2 to 13 depict amino acid sequences which are in the order of claims 2 to 13 21, **21**, **19**, **18**, **20**, **23**, **14**, **22**, **23**, **24**, **25** and **26** amino acids long) and when each of X and Y is single bond, are peptides of (or consisting of) 5 to 30 amino acids which bind to IL-6. They are stated in D5 (see, e.g., page 57, lines 35-to 38) to inhibit the binding of IL-6 to the IL-6 receptor and it is also stated there that the administration of the peptides to a patient with an autoimmune disease inhibits the production of autoimmune antibodies caused by binding of IL-6 to its receptor. The peptides therefore antagonize the activity of IL-6 in a patient with autoimmune

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disease.

Therefore, independent claim 1 (relating to peptides consisting of 5 to 30 amino acids and exhibiting antagonistic activity directed against IL-6) and independent claim 3 (relating to peptides consisting 5 to 30 amino acids and exhibiting antagonistic IL-6 activity) appear to lack novelty over D5.

- 3.2 The above said 19, 18 and 14 amino acid long peptides of D5 appear to anticipate claim 4 (relating to peptides consisting of 5 to 20 amino acids and exhibiting antagonistic activity directed against IL-6).
- 3.3 The above said 11 D5 peptides do not anticipate claim 2. D8 (which is, according to your last letter, prior art) shows that the IL-6 receptor consists at least of the alpha-chain (subunit) and of the beta-chain (subunit) (see page 714, right hand column, first full sentence). We are, however, unable to show that the above said 11 D5 peptides exhibit antagonistic activity directed against the alpha- or beta-chain of the IL-6 receptor. The peptides could bind the receptor at a different place of the receptor.
- 3.4 D5 also discloses the use of the above said 11 peptides for the production of antibodies, for therapeutical and diagnostic purposes and also for removing IL-6 from body fluids. Therefore, claims 13-24 also appear to lack novelty over D5.
- 3.5 Note that all claims which we have not objected in above items 2.1 to 2.4 are considered novel over D5.
- 3.6 Claim 12 appears not to involve an inventive step (Art. 33(3) EPC). Making a mixture of at least two of the above said 11 known D5 peptides is not considered and inventive contribution to the art.
- 4. Art. 33(2) over D2.
- 4.1 The IL-6 antagonizing decapeptide RYILDGISAL of D2 falls within claim 1 (mentioning a peptide of 5-30 amino acids), claim 3 (mentioning a peptide of 5-30 amino acids), claim 4 (mentioning a peptide of 5-20 amino acids) and

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claim 5 (mentioning a peptide of 5 to 12 amino acids). Hence, these claims appear to lack novelty.

The peptide is stated in D2 to be useful together with a carrier for treating various autoimmune diseases. Therefore, claims 15 and 16 (relating to a pharmaceutical preparation and its use) and 24 (relating to the use of the peptide for preparing a medicament) also appear to lack novelty.

- 4.2 Note that all claims which we have not objected in item 4.1 are considered novel over D2.
- Art. 33(2) and (3) EPC over D6 5.
- 5.1 The IL-6 antagonizing peptides of the formula of claim 1 of D6, wherein X1 and Y1 are absent (note that X1 and Y1 are stated in the claim to be optional), wherein X2 = hydrogen and wherein Y2 = OH (these peptides consists of 8 amino acids) anticipate claims 1, 3, 4 and 5 which therefore appear to lack novelty.
- 5.2 The IL-6 antagonizing peptides of the formula of claim 7 of D6, wherein X1 and Y1 are absent, wherein X2 = hydrogen and wherein Y2 = OH (these peptides consists of 5 amino acids) anticipate claims 1, 3, 4 and 5 which therefore appear to lack novelty.
- 5.3 D6 also discloses the use of its peptides for the production of antibodies and for therapeutical and diagnostic purposes. Therefore, it also appears that claims 13 to 16, 18, 19 and 22 to 24 lack novelty over D6.
- 5.4 Note that all claims which we have not objected in above items 5.1 to 5.3 are considered novel over D5.
- 5.5 Claim 12 does not lack novelty over D6. But, making a mixture containing at least two of the known peptides is not considered an inventive contribution to the art. Claim 12 appears not to involve an inventive step (Art. 33(3) EPC).
- The remaining documents D1, D3, D4, D7 and D8 appear not to anticipate/ 6.

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render obvious more of the claims than D2, D5 and D6 as discussed hereinbefore do.

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<u>CLAIMS</u>

- A peptide of 5-30 amino acids which peptide exhibits antagonistic activity directed against IL-6.
- A peptide of 5-30 amino acids which peptide exhibits antagonistic activity directed against the α or β -chain of the IL-6 receptor.
- A peptide of 5-30 amino acids which peptide exhibits antagonistic or agonistic IL-6 activity.
- A peptide according to claim 1, 2 or 3 of 5-20 amino acids.
- A peptide according to claim 4 of 5-12 amino acids. 10
 - A peptide according to claim 1, 2, 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences:
 - STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM QLSCFRKSPLSNVVC.
- PRSTPSLTTKAVLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNE-15 GKKMRCEWDGGR, NFTLKSEWATHKFADCKAKRDIPIS, WVEAENALGKVTSDH, or PVYKVKPNPPHNLSVIN.
 - A peptide according to claim 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with
- the following amino acid sequence: EWGPRSTPSLTT-20 KAVLLVRKFQNSPAED
 - A peptide composition, wherein at least two peptides according to any of claims 1-7 are chemically linked directly or via spacer molecules.
- A peptide composition according to claim 8 wherein at 25 least two peptides are linked with lysine.
 - A peptide composition wherein at least two peptides according to claim 1, 2, 3, 4, or 5 having at least one string of 5 consecutive amino acids long in common with the
- amino acid sequence RYILDGISALRK are linked with lysine. 30

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- 11 A peptide composition according to claim 8, 9 or 10 wherein at least four peptides are linked with branching oligolysines.
- 12 A mixture comprising peptides and/or peptide compositions according to any of claims 1-11.
- 13 Antibody specifically directed against a peptide cr a peptide composition according to any of claims 1-11.
- 14 Anti-idiotypic antibody raised against an antibody according to claim 13.
- 10 15 A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to any of the above claims together with at least one suitable excipient for administration.
 - 16 Use of a pharmaceutical preparation according to claim
- 15 15 in the treatment or prevention of an IL-6 related disease.
 - 17 Use of a peptide, peptide composition or antibody according to anyone of claims 1-13 to clear extra-corporeal blood or blood products from IL-6 or IL-6 receptor molecules.
- 18 A diagnostic assay comprising a peptide or a peptide 20 composition or an antibody according to anyone of claims 1-13.
 - 19 Use of a diagnostic assay according to claim 18 to detect or diagnose IL-6 related disease in man or animals.
 - 20 Use of a peptide according to claim 7 to exert agonistic
- 25 IL-6 activity at concentrations that are relatively equivalent to 7.5 to 120 μ g/ml when tested in vitro in a B9 cell bio assay.
 - 21 Use of a peptide according to claim 20 in cell-culture.
- 22 A pharmaceutical preparation comprising a peptide 30 according to claim 7 together with at least one suitable excipient for administration.
 - Use of a pharmaceutical preparation according to claim 22 for topical or intra-mammary application.
- 24 Use of a peptide, or peptide composition according to anyone of claims 1-12 for the manufacture of a medicament for topical or intra-mammary application.

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(54) Title: IL-6 AND IL-6-RECEPTOR DERIVED PEPTIDES HAVING IL-6 ANTAGONISTIC OR AGONISTIC ACTIVITY

(57) Abstract

The invention relates to IL-6 and IL-6-receptor derived peptides having IL-6 agonistic or antagonistic activity. The peptides are at least 5 amino acids long and are selected from one of the following amino acid sequences: RYILDGISALRK, STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQMQLSCFRKSPLSNVVC, PRSTPSLTTKAVLLVRKFQNS, MCVASSVGSKFSK-TQTFQGC, PEKPKNLSCIVNEGKKMRCEWDGGR, NFTLKSEWATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, EWGPRSTP-SLTTKAVLLVRKFQNSPAED or PVYKVKPNPPHNLSVIN. Selected peptides and combinations of selected peptides can be used in the treatment, prevention, detection, or diagnosis of IL-6 related disease and can be used to clear blood or bloodproducts of IL-6 or IL-6 receptor molecules.

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IL-6 AND IL-6-RECEPTOR DERIVED PEPTIDES HAVING IL-6 ANTAGONISTIC OR AGONISTIC ACTIVITY

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The invention relates to the field of cytokines. Cytokines are substances that are produced by cells of the immune system and are involved in regulation of humoral and cellular immune reactions and inflammatory responses. Many cytokines are known, and all exert influence on various reactions in the body in a complicated fashion. To illustrate their interdependency and the intricate web of relationships that exist between cytokines, one often speaks about the "cytokine network".

Interleukine 6 (IL-6) is a cytokine which has many effects upon mammalian cells. It exerts these effects through binding to a specific cell surface receptor, that consists of a specific α-subunit of with a molecular weight of approximately 80 kD and a common β-subunit of approximately 130 kD, also named gp130. The gp130 ß chain is also involved in signal transduction of interleukin-11 (IL-11), leukemia inhibitory factor (LIF), oncostatin M (OM), ciliairy neurotrophic factor (CNTF), and cardiotrophin-1 (CT-1) (P.B. Sehgal, Ling Wang, Ravi Rayanade et al., pp 1-14; volume 762, Annals of the New York Academy of Sciences; 1995).

IL-6 is an extremely pleiotropic cytokine, and its activities include: induction of Ig production by B cells, stimulation of B and T cell growth, differentiation of T cells and macrophages, induction of acute phase protein production by hepatocytes, multilineage hematopoesis, osteoclast formation, maturation of megakaryocytes, and platelet production. IL-6 also effects the central nervous system: IL-6 is an endogenous pyrogen and can induce ACTH production by the pituitary, finally resulting in increased glucocorticoid levels in the circulation. IL-6 exerts its activity by triggering a transmembrane receptor that is present on all target

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cells. Specific steps in the IL-6 signaling cascade are the binding to the low affinity α -chain (CD126). The complex of IL-6 and α -chain binds with the high affinity signal transducing β -chain (GP130, CD130).

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In healthy individuals no or only very low levels of IL-6 (<10 pg/ml) are detectable in the circulation. IL-6 levels are increased in various diseases, and it is postulated that these increased levels play a causative role in the pathogenesis of these diseases. Examples of diseases where increased levels of IL-6 are found are multiple myeleoma, AIDS lymphoma, polyclonal B cell activation as observed in AIDS, rheumatoid arthritis, cardiac myxoma and Castleman's disease, mesangial proliferative glomerulonephritis, psoriasis, cancerassociated cachexia, postmenopausal osteoporosis, sepsis, multiple system organ failure, alcohol cirrhosis, and diseases of the central nervous system like Alzheimer, among others. Evidence for the causative role of IL-6 in the pathogenesis of some of the above mentioned diseases has come from from phase I/II clinical trials with IL-6 neutralizing monoclonal antibodies. Treatment with anti-IL-6 monoclonal antibodies reversed fever, acute phase proteins, night sweats, bone destruction, and cachexia. Treatment of a patient with Castleman disease with anti-IL-6 monoclonal antibodies reduced acute phase protein levels, fever, anemia, thrombocytosis, and hypergammaglobulinemia. Improvement of patients was also observed in patients with rheumatoid arthritis. Apparently, reduction of IL-6 activity in these patients resulted in improvement of the clinical signs of their disease.

This approach for treating disease by antagonizing IL-6 activity makes use of monoclonal antibodies directed to IL-6. However, monoclonal antibodies are usually not of human origin and repeated administration of non-human monoclonal antibodie generally leads to immune responses against the constant part of the antibodies, since this is foreign to the body of the patient. This

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immunereaction to the monoclonal antibodies used in the treatment is, first of all, counterproductive to the therapeutic treatment itself. The monoclonal antibodies used will be rendered ineffective by the reaction with the antibodies produced by the immune system. Secondly, repeated administration of non-human monoclonal antibodies may elicit such severe immune reactions that they will be detrimental to the patient. Methods for producing less antigenic antibody fragments and methods for humanizing antibodies have been proposed, but, if feasible at all, these methods are not very economical and will their own give rise to problems regarding to half-life and bio-availability. Consequently, using anti-IL-6 monoclonal antibodies in the treatment of IL-6 related disease is considered not to be feasible.

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Inhibitors or antagonists based upon mutagenesis of IL-6 have also been proposed, such as IL-6.Q160E /T163P (Brakenhoff, J., de Hon, F., Fontaine, V., et al; J.Biol.Chem.; 269:86-93 (1994)), and IL-6.Q159E/T162P (Ehlers, M., de Hon, D., Klaasse Bos, H, et al., J. Biol. 20 Chem.; 270:8158-8163 (1995)). It has been shown with these mutant proteins that receptor binding of IL-6 and signal transduction of IL-6 can be separated in vitro. such mutant proteins are also foreign to the 25 body of the patient to be treated and will also elicit an unwanted and unfavourable immune response that generally is detrimental to the treatment. Furthermore, such mutant proteins may only be partly effective, in that, although they may effectively block or inhibit specific IL-6 activities, at the same time they may exert other effects 30 on the cytokine network with additional, still remaining, reactive sites present on these proteins. Therapeutic treatment with such reagents would then elicit other, yet unpredictable, side effects. A great disatvantage of earlier reported mutant IL-6 and IL-6 receptor 35 antagonists is that these molecules, instead of inhibiting IL-6 in vivo, act as carrier and increase the

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half-life time and result in an increase of IL-6 activity in vivo. Moreover, these mutant IL-6 and IL-6 receptor antagonists have a low affinity to their target molecules and will likely act as an immunogen. In addition, antibodies raised to IL-6 stabilize IL-6 and result in an increased IL-6 production. Accumulation of circulating IL-6 as stable IL-6-anti-IL-6 complexes as a result of treatment with these antibodies to IL-6, will occur as no renal filtration can be expected. Repeated use of nonhumanized IL-6 antibodies to human patients will most likely induce antibody production to these antibodies, and result in formation of immune-complexes (Heremans, H., Dillen, C., et al J.Immunol. 22,2395-2401).

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The present invention provides a solution to the above illustrated problems without hampering the 15 possibility of therapeutic treatment of IL-6 related disease. The above methods to inhibit IL-6 activity by antibodies or mutants, differ greatly from the invention as described here: peptides that antagonize or agonize IL-6 at the binding site to the receptor in three ways: 20 at the IL-6 part, at the α -receptor part, and at the gp130-receptor part. These antagonists and agonists and combinations of these antagonists and/or agonists as multimeric peptides or as single peptides with defined pharmacokinetic characteristics gives a powerful tool to 25 manage IL-6 bioactivity. With the solution provided by the present invention, immune responses to the treatment do not occur. Further, the occurence of unpredictable side effects is greatly minimized.

The invention provides synthetic peptides that interact with the receptor site of IL-6 or with IL-6 receptors (α and β) present at target cells.

The invention further provides synthetic peptides that, when combined, interact with the receptor site of IL-6 as well as with IL-6 receptors (α and β) present at target cells. A mixture of these peptides is particulary valuable as the pharmacological properties of the

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peptides can be adjusted to obtain a maximal desired effect. Moreover, half-life time can be prolonged by inserting unnatural amino-acids into the synthetic peptides. The antagonizing or agonizing activity of the peptides is increased by producing di- or multi-meric peptides directed to one or more receptor sites. Such dior multimeric peptides can for instance be made by linking the peptides via one or more amino-acids such as lysine (Tam, PNAS 1988, 85: 5409-5413). The distribution 10 of the peptides into target organs can be optimized by adjusting the hydrophilic or lipophylic nature of peptides or by binding of these peptides onto peptides that interact with specific organ markers. Finally, the peptides provided can be bound onto the solid phase of 15 membranes or filters that are connected into an extracorporal blood circulation circuit of the patient. A more efficient clearance of IL-6 and/or soluble IL-6 receptors can in that way be achieved.

Such synthetic peptides can be derived from (A) IL-6, or derived from (B) the receptor α -chain of IL-6 (IL-20 6Rα, CD126), or from (C) the receptor ß-chain of IL-6 (IL-6RS, GP130, CD130) and exhibit antagonistic and agonistic activity against the various components and steps of the IL-6 signaling cascade. The peptides were found by testing sets of overlapping amino acid sequences 25 from the published human IL6 (Hirano, T., Yasukawa, K., Harada, H., et al.; Nature 324, 73-76 (1986); Yasukawa, R., Hirano, T., Watanabe Y., et al.; EMBO J. 6:2939-2945 (1987), IL-6Ra (Yamasaki, K., Taga, T., Hirata, Y., et al.; Science 241:825-828 (1988)) and IL-6Rß (Hibi, M., 30 Murakami, M., Saito, M.; Cell 63:1149-1157 (1990)). These overlapping peptides, each twelve amino acids long, were tested in an assay for antagonistic or agonistic IL-6 activity.

35 The peptides provided by the invention all exhibit antagonistic or agonistic IL-6 activity against the IL-6 signaling cascade as measured in an IL-6 assay. The

peptides of the present invention are too small to generate immune responses. Further, they are too short to contain additional reactive sites, so that the antagonistic and, in addition, the agonistic peptides can advantageously be used to treat patients to counteract and adjust elevated IL-6 levels. The amino acids in all antagonistic or agonistic peptides described below are identified by the one letter code, in which the N-terminal (head) amino acid is listed first (on the left) and the C-terminal (tail) amino acid is listed last (on the right).

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A. The antagonistic peptides derived from IL-6 preferably comprise at least 5 consecutive amino acids selected from the following 3 regions that were identified as RYILDGISALRK, STKVLIQFLQKKAKNL, and I-LRSFKEFLOSSLRALROM.

- B. The antagonistic peptides derived from the receptor α -chain of IL-6 preferably comprise at least 5 consecutive amino acids selected from the following 3 regions that were identified as QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLVRKFQNS, and MCVASSVGSKFSKTQTFQGC. The agonistic peptides derived from the receptor α -chain of IL-6 preferably comprise at least 5 consecutive amino acids selected from the following region that was identified as EWGPRSTPSLTTKAVLLVRKFQNSPAED.
- C. The antagonistic peptides derived from the receptor ß-chain of IL-6 preferably comprise at least 5 consecutive amino acids selected from the following 4 regions that were identified as PEKPKNLSCIVNEGKKMRCE-WDGGR, NFTLKSEWATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, and PVYKVKPNPPHNLSVIN.

Relatively short peptides (as short as a string of 5 amino acids) that are selected from any of the above peptides, or peptides of no more than 30 amino acids long which show antagonistic or agonistic activity as measured in an IL-6 assay and have at least one string of at least 5 amino acids in common with the peptides from groups A,

B or C, are also peptides of the present invention. The peptides according to the invention can vary in length. Also, the peptides comprising a string of at least 5 amino acids which are in common with the peptides from groups A, B, and C can be modified by replacing one or a few amino acids in said string by other amino acids. Such amino acids can be selected from any of the naturally occuring amino acids, but also amino acids that normally do not occur in nature can be used as replacement amino acid. The choice of the replacing amino acid can for example be guided by comparing IL-6 or IL-6 receptor sequences from other species than humans or by selecting amino acids that lead not to extreme functional or conformational changes of the selected peptide, but also other selection methods can be used. More in particular, the present invention relates in a first aspect to a peptide containing at least 5 amino acids and at most 30 amino acids that exhibits antagonistic activity directed against IL-6 and/or against the α -chain of the IL-6 receptor and/or against the ß-chain of the IL-6 receptor.

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Also, the present invention relates in another aspect to a peptide containing at least 5 amino acids and at most 30 amino acids that can exhibit antagonistic or agonistic IL-6 activity, depending on the concentration in which it is used. An example of such peptides are peptides selected with as basis with the amino acid sequence EWGPRSTPSLTTKAVLLVRKFQNSPAED as found in the α chain of the IL-6 receptor. Surprisingly, peptides selected on the basis of the aforementioned sequence expressed antagonistic IL-6 activity at high concentrations whereas at low concentrations a marked agonistic activity was found. Agonistic activity was observed in the in vitro bioassay in a concentration range from 7.5 to 120 μ g/ml peptide. At a concentration of ≥120 μq/ml these peptides had an antagonistic effect upon the biological activity of IL-6 in the bioassay. The agonistic peptides can be used in vivo in concentrations

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that are relatively equivalent but not necessarily the same as when used in vitro.

Furthermore, the invention provides combinations of peptides, either provided as a simple mixture of several, possibly modified, peptides selected from groups A, B or C, or, provided as, possibly modified, peptides selected from groups A, B, or C that are linked, with direct chemical bonds or using spacer molecules, head to tail, or head to head, or tail to tail, or via side chains of the amino acids present in the selected peptides. 10 Examples of such combinations of peptides are for example using the peptides SLTTKAV and ILRSFKEFLQSS, or WVEAENALGKVTSDH and RYILD, or KAVLLVRK and KAVLLVRK, but many other combinations of two or more peptides can be selected from the peptides listed in groups A, B or C. 15 Such combinations of peptides, be it simple mixtures or bound peptides, can advantageously be used to counteract the events occuring in the IL-6 signaling cascade, such as disrupting the binding of IL-6 to the α -chain by 20 simultanous competing at both the IL-6 and the α -chain binding site, or simultanous competing at the binding sites of the IL-6/ α -chain complex and the ß-chain.

The peptides of the invention can suitably be used in a medicinal or pharmaceutical preparation for therapeutic or prophylactic purposes. Further, they can 25 be used in protocols to remove circulating IL-6 from the blood of diseased patients via dialysis methods in which the peptides are bound to a solid phase. Passing blood or blood filtrates along the thus bound peptides will result in clearance of IL-6 that will bind to the peptide at the 30 solid phase. Also the peptides according to the invention may be added to blood or blood filtrates and (ir) reversibly bind to IL-6 or IL-6 receptor molecules and thus render these inactive before they re-enter the body. Also, the peptides can be used in diagnostic tests, i.e. 35 in direct binding or competition based enzyme-linked immunosorbent assays to measure IL-6 levels.

IL-6 agonistic peptides can completely or partially replace IL-6 that is added to cell-cultures, for example IL-6 is used to grow or culture IL-6 dependant cells, like B-cell hybridomas to which IL-6 as growth factor is often added, bot also cell-cultures in general will benefit from the addition of agonistic IL-6 peptides. The IL-6 agonistic peptides administered to humans or animals can be used to enhance the immune response of an host exposed to a specific immunogenic substance. The IL-6 agonistic peptides can be administered to humans or animals to increase the responsiveness of the immune system of the host. A specific use is in pharmaceutical preparations for topical or intramammary application. When these agonistic peptides are combined with IL-6 antagonists as described, excess of IL-6 can be inhibited without loss of basal IL-6 signal transduction.

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Antibodies specifically directed against the peptides, and their corresponding anti-idiotypic antibodies, are part of the invention. Such antibodies can for example be administered to patients treated earlier with the peptides, to counteract the effect of the peptides on the patient. Such antibodies can be used in the above described dialysis protocols and diagnostic tests.

Synthesis of the peptides may be acomplished according to the available methods in the art. The synthesis of the exemplified peptides was done according to Valerio et al. (Int. J. Peptide Res., 42:1-9 (1993) and/or Valerio et al. (Int. J. Peptide Res., 44:148-165 (1994)). Methods for large scale production of syntethic peptides and the purification thereof are well known in the art. The invention is illustrated in the following experimental part.

WO 97/48728

Experimental part

1. Peptide synthesis.

The peptides of the examples which were intended for identifying active centers in the IL-6 and IL-6 receptor molecules were synthesized using a method according to Valerio et al. (Int. J. Peptide Res., 42:1-9 (1993) and/or Valerio et al. (Int. J. Peptide Res., 44:148-165 (1994)). Multimeric peptides (four branched) were synthetized by the solid-phase method and using of a dispersed system with branching oligolysines as a scaffolding for incorporation of the synthetized antagonistic peptides (Tam, J.P.; Proc. Natl. Acad. Sci. USA, 85:5409-5413 (1988).

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2. Proliferation assay to determine antagonistic IL-6 activity.

A set of overlapping peptides, each twelve amino acids long (each consecutive peptide shifts one amino acid, so consecutive peptides have 11 amino acids in common), derived from human IL-6 sequence (Hirano, T., Yasukawa, K., Harada, H., et al.; Nature 324, 73-76 (1986); Yasukawa, R., Hirano, T., Watanabe Y., et al.; EMBO J. 6:2939-2945 (1987), were incubated with cells (B9) at 37°c. After one hour, recombinant human IL-6 (CLB, Amsterdam, The Netherlands) was added at 3 different concentrations (2.5 U/ml, 5 U/ml and 10 U/ml).

A set of overlapping peptides, each twelve amino acids long (each consecutive peptide shifts one amino acid, so consecutive peptides have 11 amino acids in common), derived from human IL6Ra (Yamasaki, K., Taga, T., Hirata, Y., et al.; Science 241:825-828 (1988)) or gp130 (Hibi, M., Murakami, M., Saito, M.; Cell 63:1149-1157 (1990)), were incubated with 3 different concentrations IL-6 (2.5 U/ml, 5 U/ml, 10 U/ml) diluted in DMEM supplemented with HT for one hour at 37°c. Then the residual IL-6 activity was determined in a biological

assay by measuring the IL-6 dependant proliferative growth of B9 mouse hybridoma cells (Helle, M., Boeije, L., Aarden, L.A.; Eur. J. Immunol. 18:1535-1540 (1988)). Briefly, B9 mouse hybridoma cells were collected during their logarithmic growth phase in IL-6 free media and 5 suspended at a concentration of 1*10⁵ cells/ml in DMEM+HT medium containing 5% FCS. Fifty μ l of each IL-6 dilution was combined with each of the synthetized peptides representing IL-6 sequences and incubated for 1 hour at 37°C. This mixture was added in duplicate to 50 μ l of the 10 B9 cell suspension in flat-bottommed 96-well tissue culture plates (Greiner) and incubated at 37°c and 5% CO2 for 72 h. IL-6 activity was assessed by using 3-(4,5dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT, Sigma). After addition of 25 μ l of MTT (5mg/ml 15 dissolved in PBS) to each well and further incubation at 37°c for 4 h, 100 μ l of lysis buffer (20% w/v SDS in 50 % dimethyl formamide) was added. Thereafter, incubation was continued over night at 37°c and the next morning absor-20 bance was read at 578 nm. To determine the agonistic or antagonistic activity to IL-6 of the peptides synthetized from the sequences of the IL-6 receptor a or ß, various concentrations of each of these peptides was combined with 50 μ l of the B9 cell suspension (1*10⁵ cells/ml in DMEM+HT medium containing 25 5% FCS). This suspension was incubated for 1 hour at 37°C, and combined with each of the dilutions of IL-6 into flat-bottommed 96-well tissue culture plates (Greiner). Plates were incubated at 37°C for 72 h. IL-6 activity was assessed as described above. 30 Samples without synthetized peptides or with a sham peptide but with IL-6 were used as positive control, whereas samples that contained neither IL-6 nor

35 Inhibition or enhancement of IL-6 activity was determined by calculating the ratio absorbance of test sample and

synthetized peptides were used as negative control.

absorbance positive control both corrected for negative control absorban ce.

3. Toxicity testing of peptides

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Three separate test were performed to determine whether the synthetized peptides excert toxic effect in vitro upon erythrocytes (A), or polymorphonuclear cells (B), or hepatocytes (C).

- A. Sheep red blood cells (SRBC) were washed five times in PBS. A 1% (v/v) suspension of erythrocytes was prepared in veronal-buffered saline that contained gelatin (GVS: 0.032% gelatin in 3.9 mM barbitone sodium, 1 mM MgSO₄, 0.38 mM CaCl₂, and 145.6 mM NaCl). Twofold dilutions of the synthetized peptides (50 μl) were made in U-shaped microtiter plates (Greiner Labortechnik) and 50 μl of the SRBC suspension were added to each well. Plates were sealed, mixed and incubated for 2 hours at 37°C. Thereafter, plates were examined for hemolysis. None of the synthetized peptides showed hemolysis.
- B. Porcine polymorphonuclear cells (PMN) were isolated from pig blood (Cruijsen, T.L.M., Van Leengoed, L.A.M.G. et al.; Infect. Immun. 60:4867-4871 (1992)). Twofold dilutions of the synthetized peptides (50 μ l) were made in flat-bottomed microtiter plates (Greiner Labortechnik)
- and 50 μ l of the PMN suspension (2*10⁶ cells/ml) were added to each well. Plates were sealed, gently mixed and incubated for 6 hours at 37° and 5% CO₂. Thereafter, plates were examined for cytotoxicity by nigrosine dye exclusion. Non of the synthetized peptides was toxic for PMN.
 - C. Porcine hepatocytes were isolated from liver of pigs based on Seglens' method (Seglen, P.O.; Methods Cell Biol 13:29-83 (1976)) and adapted according to Monshouwer M., et al. (Toxicol. Applied Pharmacol.in press). Hepatocytes were suspended in Williams' medium E to a concentration of 10⁶ cells/ml. From this suspension 1.5 ml was put into each well of 12-well tissue culture plates (Costar) and

incubated for 12 h at 37°C. Adherent hepatocytes were examined for their viability and nonadherent hepatocytes were discarded. Each synthetized peptide was mixed with Williams' medium E (at dilutions of 1:50 and 1:100) and added to wells with adherent hepatocytes. After another 24 h incubation at 37°C viability was assessed by using 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT, Sigma). After addition of 1.5 of MTT (1-mg/ml dissolved in Williams' medium E) to each well and further incubation at 37°c for 30 min, 1 ml of lysis buffer (0.8 M HCL in isopropanol) was added. Thereafter, plates were mixed for 10 min and absorbance was read at 560 nm. None of the synthetized peptides proved to affect the viability of the hepatocytes.

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4. Effect of IL-6 antagonistic peptides upon IL-6 induced acute phase reaction and downregulation of hepatic biotransformation activities.

Porcine hepatocytes were isolated from liver of pigs based on Seglens' method (Seglen, P.O.; Methods Cell Biol 20 13:29-83 (1976)) and adapted according to Monshouwer M., et al. (Toxicol. Applied Pharmacol.in press). Hepatocytes were suspended in Williams' medium E to a concentration of 106 cells/ml. From this suspension 1 ml was put into each well of 12-well tissue culture plates (Costar) and 25 incubated for 12 h at 37°C. Adherent hepatocytes were examined for their viability and nonadherent hepatocytes were discarded. Each synthetized peptide was mixed (at dilutions of 1:50 and 1:100) with Williams' medium E containing IL-6 (1000 U/ml) and added to wells with 30 adherent hepatocytes. Also a negative (containing no IL-6 and without synthetized peptides in the medium) and positive control (containing 1000 U/ml IL-6 in the medium) were prepared and tested. After an incubation period of 24 hours, the medium was removed and for each 35 well CYP450 dependent enzyme activity of intact monolayers of hepatocytes was determined.

CYP450 enzym assay. CYP450 dependent enzym activity, using testosterone (250 μ M) as substrate, was determined as previously described by Van 't Klooster et al.(Bioch. Pharmacol.46;1781-1790 (1993)). Briefly, testosteron was mixed with Williams' medium E without fetal calf serum and added to the wells with hepatocytes. After 30 min incubation at 37°C and 5%CO₂, hydroxylated testosteron metabolites in the medium were quantified by HPLC.

HPLC analysis. Aliquots of 1 ml of medium was mixed with 100 μ l of a solution of 11ß-testosteron (12,5 μ g/ml) 10 in methanol as internal standard and extracted with 5 ml dichlormethane. The organic phase was transferred to clean tubes and evaporated to dryness at roomtemperature under a stream of nitrogen. The residues were dissolved in 130 μ l 50% methanol and 20 μ l of these solutions were 15 injected for HPLC analysis. The stationary phase consisted of a C18 glasscolumn (20 cm, 3 µm particle size, Chrompack, Middelburg, the Netherlands). The mobile phase consisted of buffer A (12% methanol, 75% milli Q water) and buffer B (64% methanol, 6% acetonitril, 30% milli Q 20 water). With these buffers an elution gradient was generated; 10-58% B from 0-45 minutes; 58-59% from 45-50 minutes; 59-10% from 50-53 minutes, with a flow rate of 0,8 ml/min. Metabolites were detected spectrofotometrically at 254 nm. Inhibition of IL-6 25 dependant downregulation of cytochrome P450 was determined by comparing the relative concentration of hydroxylated testosteron metabolites in medium from adherent hepatocytes incubated with synthetized peptides and IL-6, and the relative concentration of hydroxylated 30 testosteron metabolites in medium from positive and

5. Results

Peptides derived from hIL-6, hgp130 (the ß-chain of the IL-6 receptor) and hIL6Ra (the α-chain of the IL-6 receptor) were analysed for antagonistic IL-6 activity.

negative control hepatocyte monolayers.

For hIL-6 peptides, 3 regions were identified which inhibited IL-6 activity in an IL-6 assay (fig. 2). Peptide 31, 119-123 and 167-174 represent the identified regions (RYILDGISALRK, resp. STKVLIQFLQKKAKNL, resp. ILRSFKEFLQSSLRALRQM).

For hIL6Ra, also 3 regions were identified which inhibited IL-6 activity in an IL-6 asay (fig. 3). Peptide 6-9, 24-33 and 80-89 represent the identified regions (QLSCFRKSPLSNVVC, resp. PRSTPSLTTKAVLLVRKFQNS, resp. MCVASSVGSKFSKTQTFQGC).

For hgp130 peptides, 4 regions were identified which inhibited IL-6 activity in an IL-6 assay (fig. 4). Peptide 2-15, 33-46, 73-76 and 92-97 represent the identified regions

15 (PEKPKNLSCIVNEGKKMRCEWDGGR, resp. NFTLKSE-WATHKFADCKAKRDTPTS, resp. WVEAENALGKVTSDH, resp. PVYKVKPNPPHNLSVIN).

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The identified peptides with anti-IL-6 activity were not lytic for erythrocytes and not toxic for polymorphonuclear cells and not toxic for primary hepatocyte culture cells.

Peptides derived from hIL6Ra (the α -chain of the IL-6 receptor) were analysed for agonistic IL-6 activity and 1 region was identified which stimulated proliferation of B9 cells without IL-6 added to the medium and enhanced IL-6 activity in the B9 bio-assay (fig. 5). Peptide 21-37 represent the region EWGPRSTPSLTTKAVLLVRKFQNSPAED of the IL-6Ra sequence

Agonistic activity was observed in a concentration range from 7.5 to 120 μ g/ml peptide. These peptides induced prolife rative growth of the IL-6 dependant cell line B9, and when combined with IL-6 enhanced proliferation of the B9 cell line was examined, and thus the biological activity of IL-6 was enhanced. At a concentration of $\ge 120~\mu$ g/ml these agonistic peptides had an antagonistic effect upon the biological activity of IL-6.

The identified peptides with agonistic IL-6 activity were not lytic for erythrocytes and not toxic for polymorphonuclear cells and not toxic for primary hepatocyte culture cells.

Synthetized peptides from the regions PVYKVKPNPP-HNLSVIN, WVEAENALGKVTSDH, and MCVASSVGSKFSKTQTFQGC inhibit IL-6 regulated downregulation of cytochrome P-450 of hepatocytes.

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CLAIMS

- 1 A peptide containing 5-30 amino acids which peptide exhibits antagonistic activity directed against IL-6.
- 2 A peptide containing 5-30 amino acids which peptide exhibits antagonistic activity directed against the α and/or β -chain of the IL-6 receptor.
- 3 A peptide containing 5-30 amino acids which peptide exhibits antagonistic and/or agonistic IL-6 activity.
- A peptide according to claim 1, 2 or 3 containing 5-20 amino acids.
- 10 5 A peptide according to claim 4 containing 5-12 amino acids.
 - A peptide according to claim 1, 2, 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences:
- 15 RYILDGISALRK, STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM
 QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLVRKFQNS,
 MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNEGKKMRCEWDGGR, NFTLKSEWATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or
 PVYKVKPNPPHNLSVIN.
- 7 A peptide according to claim 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence:

 EWGPRSTPSLTTKAVLLVRKFQNSPAED
- 8 A peptide composition, wherein at least two peptides 25 according to any of claims 1-7 are chemically linked directly or via spacer molecules.
 - 9 A peptide composition according to claim 8 wherein at least two peptides are linked with lysine.
- 10 A peptide composition according to claim 8 wherein 30 at least four peptides are linked with branching oligolysines.

- 11 A mixture comprising peptides and/or peptide compositions according to any of claims 1-10.
- 12 Antibody specifically directed against a peptide or a peptide composition according to any of claims 1-10.
- 5 13 Anti-idiotypic antibody raised against an antibody according to claim 12.
 - 14 A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to any of the above claims together with at least one suitable excipient for administration.
 - 15 Use of a pharmaceutical preparation according to claim 14 in the treatment or prevention of an IL-6 related disease.

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- 16 Use of a peptide, peptide composition or antibody
 15 according to anyone of claims 1-12 to clear extracorporeal blood or blood products from IL-6 or IL-6
 receptor molecules.
 - 17 A diagnostic assay comprising a peptide or a peptide composition or an antibody according to anyone of claims 1-12.
 - 18 Use of a diagnostic assay according to claim 17 to detect or diagnose IL-6 related disease in man or animals.
- 19 Use of a peptide according to claim 7 to exert agonistic IL-6 activity at concentrations that are relatively equivalent to 7.5 to 120 μ g/ml.
 - 20 Use of a peptide according to claim 19 in cell-culture.
- 21 A pharmaceutical preparation comprising a peptide 30 according to claim 7 or 19 together with at least one suitable excipient for administration.
 - 22 Use of a pharmaceutical preparation according to claim 21 for topical or intra-mammary application.
- Use of a peptide, or peptide composition according to anyone of claims 1-11 for the manufacture of a medicament for topical or intra-mammary application.

Fig. 1. Amino acid sequences and sources of selected peptides.

Peptides were selected from published sequences of IL-6 (A) and IL-6 a- (B) and ß-receptor (C).

A) human IL6 (Hirano, T., Yasukawa, K., Harada, H., et al.; Nature 324, 73-76 (1986); Yasukawa, R., Hirano, T., Watanabe Y., et al.; EMBO J. 6:2939-2945 (1987)
Amino acid sequence:

APPVPPGEDSKDVAAPHRQPLTSSERIDKQIRYILDGISALRKETCNKSNMCESSK-EALAENNLNLPKMAEKDGCFQSGFNEETCLVKIITGLLEFEVYLEYLQNRFESSEEQ-ARAVQMSTKVLIQFLQKKAKNLDAITTPDPTTNASLLTKLQAQNQWLQDMTTH-LILIRSFKEFLQSSLRALRQM

B) the receptor a-chain of IL-6 (IL-6Ra, CD126), (Yama-saki, K., Taga, T., Hirata, Y., et al.; Science 241:825-828 (1988))

Amino acid sequence:

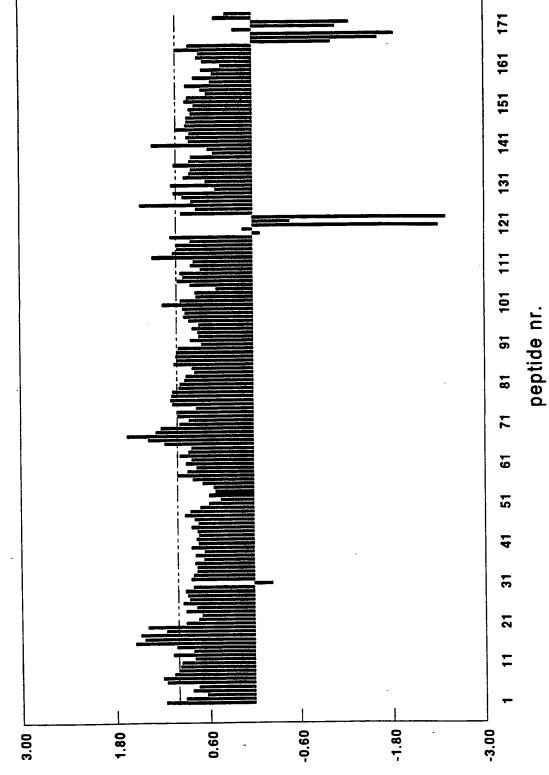
PPEEPQLSCFRKSPLSNVVCEWGPRSTPSLTTKAVLLVRKFQNSPAEDFQEPCQY-SQESOKFSCOLAVPEGDSSFYIVSMCVASSVGSKFSKTQTFQGCGILQPDPPANITV

C) the receptor &-chain of IL-6 (IL-6R&, GP130, CD130) (Hibi, M., Murakami, M., Saito, M.; Cell 63:1149-1157 (1990))

Amino acid sequence:

PPEKPKNLSCIVNEGKKMRCEWDGGRETHLETNFTLKSEWATHKFADCKAKRDT-PTSCTVDYSTVYFVNIEVWVEAENALGKVTSDHINFDPVYKVKPNPPHNLSVIN

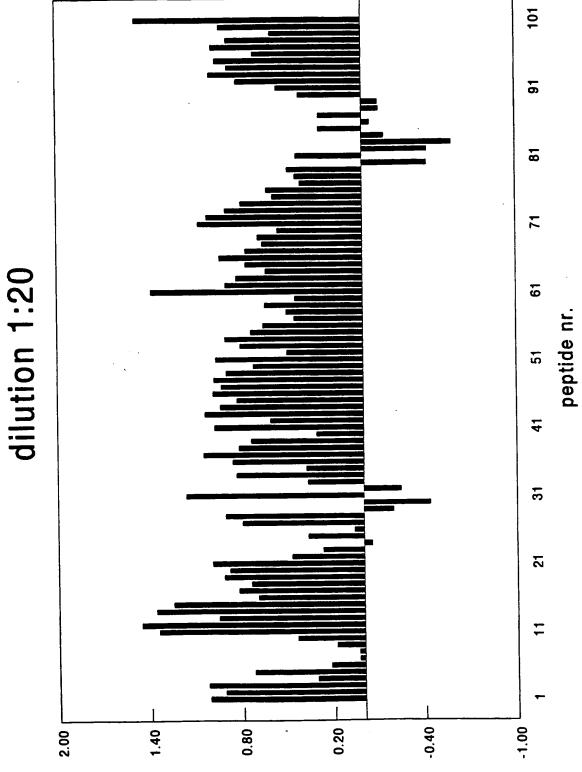




Ratio (absx - absnc)/(abspc - absnc)

Fig 2: Screening of synthetized peptides, representing IL-6, in the B9 bio-assay

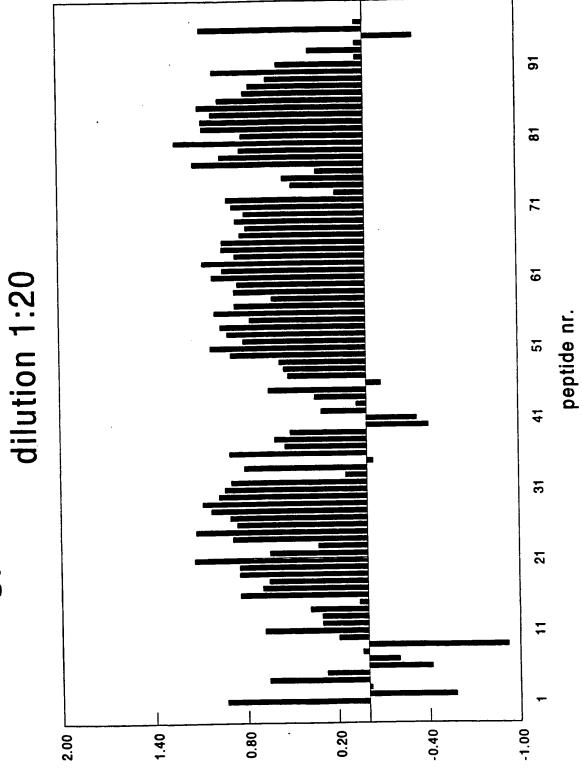




ratio (abs x-abs nc)/abs pc-abs nc)

ning of synth tiz d peptides, repr s nting IL-6Ra, in th B9 bio-assay Fig 3: Scr

gp130 receptor peptides



ratio (abs x-abs nc)/abs pc-abs nc)

Fig 4: Scre ning of synth tiz d peptides, r presenting gp130, in the B9 bioassay

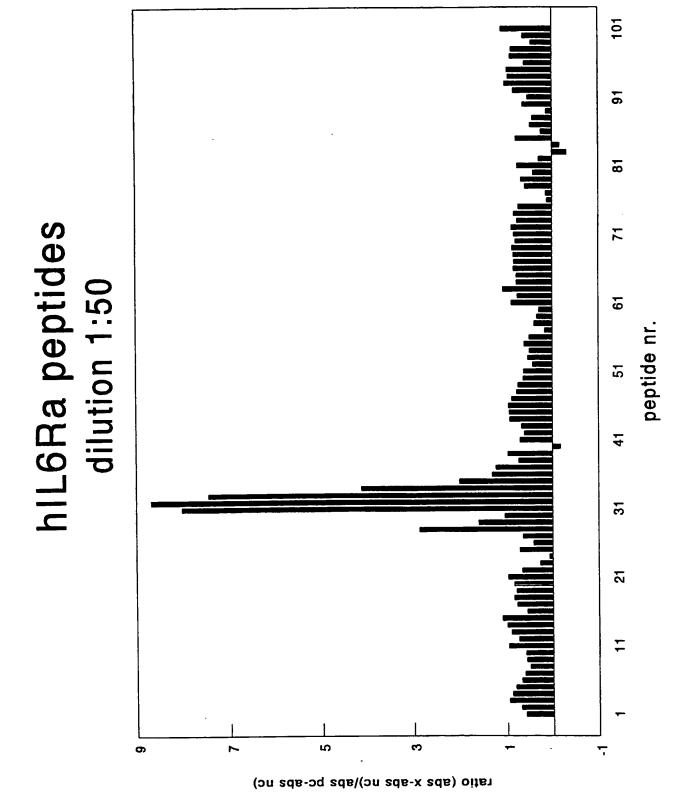


Fig 5: Screening synthetic peptides showing a region with agonistic activity (dilution 1:50)

PATENT COOPERATION TREATY

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PCT

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

Applicant's or	agent's	file reference	FOR FURTHER ACTION		Notification of Transmittal of International liminary Examination Report (PCT/IPEA/416)
PCT 0589					· · · · · · · · · · · · · · · · · · ·
nternational	applicati	on No.	International filing date (day/month)	'year)	Priority date (day/month/year)
CT/NL97	00345	5	19/06/1997		20/06/1996
nternational	Patent C	classification (IPC) or	national classification and IPC		
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pplicant					
OSTER,	Henk '	Wilhelmus et al.			
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l. This int	ernatio	nal preliminary exa	mination report has been prepare	d by this In	ternational Preliminary Examining Authorit
and is t	ransmi	tted to the applican	t according to Article 36.		
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2. This Rt	PORT	consists of a total	of 7 sheets, including this cover	sneet.	
⊠ Th	is repo	nt is also accompa	nied by ANNEXES, i.e., sheets of	the descrip	otion, claims and/or drawings
\A/I	ich ha	ve been amended.	and are the basis for this report ar	id/or sheet	s containing rectifications made
be	tore th	is Authority (see Hi	ule 70.16 and Section 607 of the A	ummstrati	ve instructions under the FOT).
Thoso	annava	es consist of a total	of 2 sheets		
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3. This re	port co	ntains indications r	elating to the following items:		
1	×	Basis of the report	•		
11		Priority			
111		•	t of opinion with regard to novelty,	inventive s	step and industrial applicability
IV		Lack of unity of inv			
V	Ø	Reasoned stateme	ent under Article 35(2) with regard	to novelty,	inventive step or industrial applicability;
		citations and expla	anations supporting such statemer	nt	
VI		Certain documents	s cited		
VII		Certain defects in	the international application		
VIII	\boxtimes	Certain observation	ns on the international application		
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		address of the IPEA/		rized officer	

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/NL97/00345

I.	Ba	sis	of	the	re	po	rt
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1. This report has been drawn on the basis of (substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments.):

		•					
	Des	cription, pages:					
	1-16	3	as originally filed				
	Clai	ms, No.:					
	1-24	ı	as received on		18/06/1998	with letter of	18/06/1998
	Dra	wings, sheets:		•			
	1-5		as originally filed				
2.	The	amendments have	e resulted in the ca	ancellation of:			
		the description,	pages:				
	\boxtimes	the claims,	Nos.:	1-23			
		the drawings,	sheets:				
3.		This report has be considered to go	een established as beyond the disclo	s if (some of) t sure as filed (l	he amendmei Rule 70.2(c)):	nts had not been m	ade, since they have been
4.	Ado	litional observation	ns, if necessary:				

INTERNATIONAL PRELIMINARY **EXAMINATION REPORT**

International application No. PCT/NL97/00345

V. R asoned stat ment under Article 35(2) with r gard to novelty, inv ntive st p or industrial applicability; citations and explanations supporting such statement

1. Statement

Novelty (N)

Yes:

Claims CLAIMS 2, 6-12

No:

Claims CLAIMS 1, 3-5, 13-24

Inventive step (IS)

Yes:

Claims CLAIMS 2, 6-11

No:

Claims CLAIM 12

Industrial applicability (IA)

Yes:

Claims CLAIMS 1-24

Claims No:

2. Citations and explanations

see separate sheet

VIII. Certain observations on the international application

The following observations on the clarity of the claims, description, and drawings or on the question whether the claims are fully supported by the description, are made:

see separate sheet

- 1.1 The documents cited in the ISR are labelled D1 to D8.
- 1.2 The claims are allowable under Art.34(1)(b) PCT.

AD SECTION VIII:

- 2. Art. 6 EPC (clarity)
- 2.1 The words " long " in **claims 6, 7 and 10** render the claims unclear, for obvious reasons.
- 2.2 The term in **claim 20** " at concentrations relatively equivalent to bio assay " lacks clarity (**Art. 6 PCT**). The clear limits of this let us say range feature are not stated in the claim and different views of the limits of the range are conceivable for different skilled persons. The claim is subject to subjective interpretation and it therefore lacks clarity.

AD SECTION V.2.:

- 3. Art. 33(2) and (3) over D5
- The peptides of the general formula of claim 1 of **D5**, when A is a peptide fragment of the formulas of claims 2 to 13 (the formulas in claims 2 to 13 depict amino acid sequences which are in the order of claims 2 to 13 21, **21**, **19**, **18**, **20**, **23**, **14**, **22**, **23**, **24**, **25** and **26** amino acids long) and when each of X and Y is single bond, are peptides of (or consisting of) 5 to 30 amino acids which bind to IL-6. They are stated in D5 (see, e.g., page 57, lines 35-to 38) to inhibit the binding of IL-6 to the IL-6 receptor and it is also stated there that the administration of the peptides to a patient with an autoimmune disease inhibits the production of autoimmune antibodies caused by binding of IL-6 to its receptor. The peptides therefore antagonize the activity of IL-6 in a patient with autoimmune

disease.

Therefore, independent claim 1 (relating to peptides consisting of 5 to 30 amino acids and exhibiting antagonistic activity directed against IL-6) and independent claim 3 (relating to peptides consisting 5 to 30 amino acids and exhibiting antagonistic IL-6 activity) appear to lack novelty over D5.

- 3.2 The above said **19, 18 and 14 amino acid** long peptides of **D5** appear to anticipate **claim 4** (relating to peptides consisting of **5 to 20 amino acids** and exhibiting antagonistic activity directed against IL-6).
- 3.3 The above said 11 D5 peptides **do not anticipate claim 2**. D8 (which is, according to your last letter, prior art) shows that the IL-6 receptor consists **at least** of the alpha-chain (subunit) and of the beta-chain (subunit) (see page 714, right hand column, first full sentence). We are, however, unable to show that the above said 11 D5 peptides exhibit antagonistic activity directed against the alpha- or beta-chain of the IL-6 receptor. The peptides could bind the receptor at a different place of the receptor.
- 3.4 D5 also discloses the use of the above said 11 peptides for the production of antibodies, for therapeutical and diagnostic purposes and also for removing IL-6 from body fluids. Therefore, **claims 13-24** also appear to lack novelty over D5.
- 3.5 Note that all claims which we have not objected in above items 2.1 to 2.4 are considered novel over D5.
- 3.6 Claim 12 appears not to involve an inventive step (Art. 33(3) EPC). Making a mixture of at least two of the above said 11 known D5 peptides is not considered and inventive contribution to the art.
- 4. Art. 33(2) over D2.
- 4.1 The IL-6 antagonizing decapeptide RYILDGISAL of D2 falls within claim 1 (mentioning a peptide of 5-30 amino acids), claim 3 (mentioning a peptide of 5-30 amino acids), claim 4 (mentioning a peptide of 5-20 amino acids) and

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claim 5 (mentioning a peptide of 5 to 12 amino acids). Hence, these claims appear to lack novelty.

The peptide is stated in D2 to be useful together with a carrier for treating various autoimmune diseases. Therefore, **claims 15 and 16** (relating to a pharmaceutical preparation and its use) and **24** (relating to the use of the peptide for preparing a medicament) also appear to lack novelty.

- 4.2 Note that all claims which we have not objected in item 4.1 are considered novel over D2.
- 5. Art. 33(2) and (3) EPC over D6
- The IL-6 antagonizing peptides of the formula of claim 1 of **D6**, wherein X1 and Y1 are absent (note that X1 and Y1 are stated in the claim to be optional), wherein X2 = hydrogen and wherein Y2 = OH (these peptides consists of **8 amino acids**) anticipate **claims 1, 3, 4 and 5** which therefore appear to lack novelty.
- 5.2 The IL-6 antagonizing peptides of the formula of claim 7 of **D6**, wherein X1 and Y1 are absent, wherein X2 = hydrogen and wherein Y2 = OH (these peptides consists of **5 amino acids**) anticipate **claims 1, 3, 4 and 5** which therefore appear to lack novelty.
- 5.3 D6 also discloses the use of its peptides for the production of antibodies and for therapeutical and diagnostic purposes. Therefore, it also appears that claims 13 to 16, 18, 19 and 22 to 24 lack novelty over D6.
- 5.4 Note that all claims which we have not objected in above items 5.1 to 5.3 are considered novel over D5.
- 5.5 Claim 12 does not lack novelty over D6. But, making a mixture containing at least two of the known peptides is not considered an inventive contribution to the art. Claim 12 appears not to involve an inventive step (Art. 33(3) EPC).
- 6. The remaining documents D1, D3, D4, D7 and D8 appear not to anticipate/

INTERNATIONAL PRELIMINARY

International application No. PCT/NL97/00345

EXAMINATION REPORT - SEPARATE SHEET

render obvious more of the claims than D2, D5 and D6 as discussed hereinbefore do.

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CLAIMS

- A peptide containing 5-30 amino acids which peptide exhibits antagonistic activity directed against IL-6.
- A peptide containing 5-30 amino acids which peptide exhibits antagonistic activity directed against the α and/or β -chain of the IL-6 receptor.
- A peptide containing 5-30 amino acids which peptide 3 exhibits antagonistic and/or agonistic IL-6 activity.
- A peptide according to claim 1, 2 or 3 containing 5-20 amino acids.
- 10 A peptide according to claim 4 containing 5-12 amino acids.
 - A peptide according to claim 1, 2, 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences:
- RYILDGISALRK, STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM 15 QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNEGKKMRCEWDGGR, NFTLKS-EWATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or PVYKVKPNPPHNLSVIN.
- A peptide according to claim 3, 4 or 5 having at 20 least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLVRKFQNSPAED
- A peptide composition, wherein at least two peptides according to any of claims 1-7 are chemically linked 25 directly or via spacer molecules.
 - A peptide composition according to claim 8 wherein at least two peptides are linked with lysine.
- A peptide composition according to claim 8 wherein at least four peptides are linked with branching 30 oligolysines.

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A mixture comprising peptides and/or peptide 11 compositions according to any of claims 1-10.

- Antibody specifically directed against a peptide or 12 a peptide composition according to any of claims 1-10.
- Anti-idiotypic antibody raised against an antibody 5 according to claim 12.
 - A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to any of the above claims together with at least one suitable excipient for administration.
- Use of a pharmaceutical preparation according to claim 14 in the treatment or prevention of an IL-6 related disease.
- Use of a peptide, peptide composition or antibody 16 according to anyone of claims 1-12 to clear extra-15 corporeal blood or blood products from IL-6 or IL-6 receptor molecules.
 - A diagnostic assay comprising a peptide or a peptide composition or an antibody according to anyone of claims 1-12.
 - Use of a diagnostic assay according to claim 17 to 18 detect or diagnose IL-6 related disease in man or animals.
- Use of a peptide according to claim 7 to exert 25 agonistic IL-6 activity at concentrations that are relatively equivalent to 7.5 to 120 μ g/ml.
 - Use of a peptide according to claim 19 in cell-20 culture.
- A pharmaceutical preparation comprising a peptide according to claim 7 or 19 together with at least one 30 suitable excipient for administration.
 - Use of a pharmaceutical preparation according to 22 claim 21 for topical or intra-mammary application.
 - Use of a peptide, or peptide composition according 23
- to anyone of claims 1-11 for the manufacture of a 35 medicament for topical or intra-mammary application.

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NATIONAL SEARCH REPORT

tion No International Ap PCT/NL 97/00345

A. CLASSIFICATION OF SUBJECT MATTER IPC 6 C07K14/54 C07K14/715 C07K16/42 C07K16/28 · C07K16/24 G01N33/577 A61K38/20 A61K39/395 G01N33/68 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols) IPC 6 C07K A61K G01N

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	DATABASE WPI Week 9607 Derwent Publications Ltd., London, GB; AN 96-065477 XP002027134 & JP 07 324 098 A (DAICEL CHEM. IND. LTD. ET AL.), 12 December 1995 see abstract	1,3,6, 14-16
x	DATABASE WPI Week 9607 Derwent Publications Ltd., London, GB; AN 96-065476 XP002027135 & JP 07 324 097 A (DAICEL CHEM. IND. LTD. ET AL.), 12 December 1995 see abstract	1-6, 14-16

 Special categories of cited documents: "A" document defining the general state of the art which is not considered to be of particular relevance 	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"E" earlier document but published on or after the international filing date "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "O" document referring to an oral disclosure, use, exhibition or other means "P" document published prior to the international filing date but later than the priority date claimed	"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art. "&" document member of the same patent family
Date of the actual completion of the international search	Date of mailing of the international search report
20 October 1997	1 0. 11. 97
Name and mailing address of the ISA European Patent Office, P.B. 5818 Patentiaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016	Authorized officer Nooij, F

Form PCT/ISA/210 (second sheet) (July 1992)

X Further documents are listed in the continuation of box C.

Patent family members are listed in annex.

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International At ...tion No PCT/NL 97/00345

Relevant to claim No.
Helevant to claim No.
1,3,6, 14-18
1-7, 14-16,21
2,3,6,7, 16
1-5,12, 14-16
1,3,4,6,
12

Form PCT/ISA/210 (continuation of second sheet) (July 1992)

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INTERNATIONAL SEARCH REPORT

Internatic pplication No. PC1/NL 97/00345

Boxi	Observations where certain claims were found unsear hable (Continuation of it m 1 of first sheet)
This Inte	ernational Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. X	Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
	see FURTHER INFORMATION sheet PCT/ISA/210
2.	Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:
з. 🗀	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
BxII	Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This Inte	emational Searching Authority found multiple inventions in this international application, as follows:
1.	As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2.	As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
з. 🗀	As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:
4.	No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark	on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International At Ition No PCT/NL 97/00345

Information on patent family members

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 9504075 A	09-02-95	AU 7341494 A CA 2168261 A EP 0715633 A JP 9501154 T	28-02-95 09-02-95 12-06-96 04-02-97
EP 426857 A	15-05-91	CA 2026881 A WO 9009396 A US 5171837 A	09-08-90 23-08-90 15-12-92
US 5210075 A	11-05-93	NONE	

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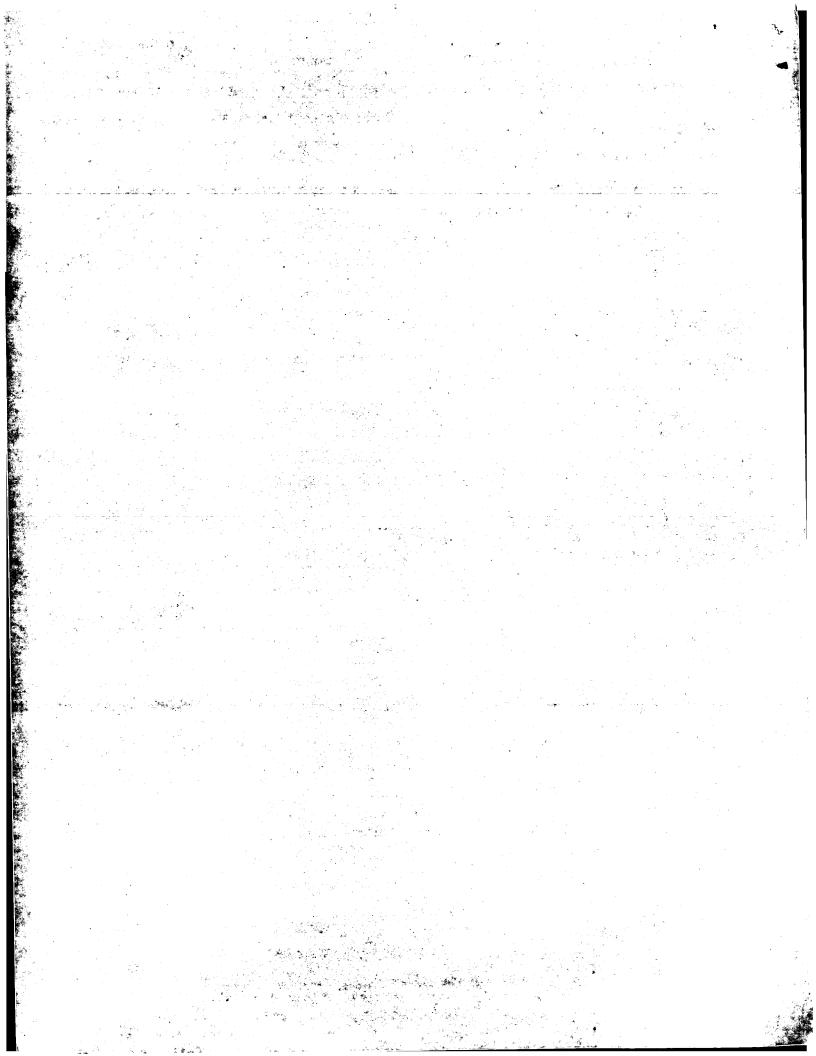
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With international search report.

(54) Title: HAEMOPOIETIC GROWTH FACTOR ANTAGONISTS

(57) Abstract

The present invention relates to modified and variant forms of haemopoietic growth factors (HGF) capable of acting as antagonists to the corresponding native haemopoietic growth factors and their use in ameliorating aberrant effects caused by the native molecules. A modified haemopoietic growth factor (HGF) is characterized by being in unglycosylated form and comprising a sequence of amino acids within a first α -helix wherein one or more exposed amino acids in said first α -helix having acidic or acidic-like properties are substituted with a basic amino acid residue. The preferred HGF are granulocyte-macrophage colony-stimulating factor (GM-CSF), interleuking (IL)-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-9, IL-10, G-CSF and erythropoietin (EPO).



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HAEMOPOIETIC GROWTH FACTOR ANTAGONISTS

The present invention relates to modified and variant forms of haemopoietic growth factors capable of acting as antagonists to the corresponding native haemopoietic growth factors and their use in ameliorating aberrant effects caused by the native molecules.

Bibliographic details of the publications numerically referred to in this specification are collected at the end of the description. Sequence Identity Numbers (SEQ ID NOs.) for the nucleotide and amino acid sequences referred to in the specification are defined following the bibliography.

Throughout this specification, unless the context requires otherwise, the word "comprise", or variations such as "comprises" or "comprising", will be understood to imply the inclusion of a stated element or integer or group of elements or integers but not the exclusion of any other element or integer or group of elements or integers.

Granulocyte-macrophage colony-stimulating factor (GM-CSF) is one member of a family of haemopoietic growth factors (HGFs) which have a similar predicted tertiary configuration (Parry et al, 1988) and whose receptors also belong to a common family (Gearing et al, 1989, Bazan, 1990). This family of haemopoietic growth factors includes, for example, in addition to GM-CSF, the cytokines IL-2, IL-3, IL-5, IL-6 and IL-10. A distinct subfamily comprising GM-CSF, IL-3 and IL-5 can be discerned based on structural similarities (Goodall et al, 1993) and on their ability to interact with a common receptor component (Lopez et al, 1992).

Human GM-CSF (hGM-CSF) comprises 127 amino acids and is available in recombinant form (rhGM-CSF). The hGM-CSF receptor has also been cloned and shown to comprise a binding (α) chain exhibiting low affinity binding to GM-CSF (Gearing *et al*, 1989) and a second (β) chain which does not measurably bind GM-CSF by itself but it allows the formation of a high affinity receptor when co-expressed with the α chain (Hayashide *et al*, 1990).

GM-CSF exhibits a range of activities extending over neutrophil, eosinophil and monocyte lineages. Specifically, GM-CSF stimulates the progenitors of these cells to proliferate and differentiate to become mature cells. In addition, it stimulates mature cells to greater function. The stimulation of mature cells results in greater capacity to phagocytose and kill micro-organisms, kill antibody-coated tumour cells and parasites and generate superoxide anion (O_2^-) in response to various stimuli. The purpose of this activation is presumed to enable the mature cells to become better effector cells in inflammatory reactions.

Therapeutically, the HGFs form an important group of molecules for their actual or potential properties. For example, the main indications for GM-CSF are for its effects on progenitor cells or mature cells. Using its effects on progenitor cells, GM-CSF is used in the treatment of bone marrow failure as seen in aplastic anaemia or chemotherapy or AIDS-induced marrow suppression. In the treatment of infections, the capacity to stimulate mature cells is especially relevant. The capacity of GM-CSF-activated neutrophils and eosinophils to kill tumour cells that have bound antibody is especially remarkable and could be used in tumour therapy.

However, despite the actual and potential benefits of HGFs, they can exhibit some detrimental side effects. For example, GM-CSF can exhibit toxicity due to stimulation of mature cells causing blood vessel damage or thrombosis. The eosinophilia caused by GM-CSF appears especially damaging in this regard. The molecule can also have detrimental effects by stimulating growth of leukaemia cells and tumour cells of non-haemopoietic origin and stimulating production of inflammatory mediators.

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International Patent Application No. PCT/AU89/00177 and an article by Lopez et al. (1992) disclose amino acid variants of GM-CSF which have exhibited reduced potency. These variants were investigated further for their potential as GM-CSF antagonists. However, the variants cause classical stimulation at concentrations 100 fold greater compared to the native GM-CSF molecule. Furthermore, attempts to find antagonistic properties failed since mixing large concentrations of one of the variants with suboptimal concentrations of native GM-CSF resulted in stronger GM-CSF stimulation with no

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evidence of inhibition being observed.

There is a need, therefore, to develop antagonists to HGFs and in particular GM-CSF which are capable of ameliorating the aberrant effects of the corresponding native molecules. There is also a need for such antagonists not to exhibit agonist properties in respect of the corresponding HGFs.

Accordingly, one aspect of the present invention provides a haemopoietic growth factor characterised by being in unglycosylated form and comprising a sequence of amino acids within a first α -helix wherein one or more exposed amino acids in said first α -helix having acidic or acidic-like properties are substituted with a basic amino acid residue.

In accordance with the present invention, it is proposed that the modified HGFs defined above act as antagonists of the native form of the corresponding HGF but not other HGFs. The term "modified" is considered herein synonymous with terms such as "variant", "derivative" and "mutant".

The HGFs are preferably GM-CSF, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-9, IL-10, granulocyte colony-stimulating factor (G-CSF) and erythropoietin (EPO) modified in accordance with the present invention. Most preferably, the HGF is GM-CSF. The HGFs are preferably in recombinant or synthetic form and, with the exception of the amino acid substitution(s) in the first α-helix, the amino acid sequence of the HGF may be the same as the naturally occurring molecule (i.e. native molecule) or may carry single or multiple amino acid substitutions, deletions and/or additions to the native amino acid sequence. The HGF sequences are preferably of mammalian origin such as from humans, livestock animals, companion animals or laboratory test animals. Most preferably, the HGFs are of human origin or of a mammalian origin capable of functioning in humans.

The first α-helix of GM-CSF has been determined at 2.4 angstrom resolution by X-ray crystallography and encompasses amino acid residues 13 to 28. Similar procedures may be adopted to determine the first α-helix in other haemopoietic growth factors. The position may also be determined by analogy to GM-CSF structure.

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Reference to "unglycosylated form" herein means that the molecule is completely unglycosylated such as when expressed in recombinant form in a prokaryotic organism (e.g. E. coli). Alternatively, a glycosylation-deficient mammalian cell may be used or complete deglycosylation may occur in vitro using appropriate enzymes. Accordingly, the present invention extends to chemically synthesised GM-CSF which is in unglycosylated form.

An "exposed" amino acid is taken herein to refer to an amino acid on an exposed or outer portion of an α -helix compared to those amino acids orientated towards the inside of the molecule.

An acidic amino acid includes, for example, Glu and Asp. Preferred basic amino acids are Arg and Lys.

- According to another aspect of the present invention, there is provided a haemopoietic growth factor characterised by:
 - (i) being in unglycosylated form;
 - (ii) comprising a sequence of amino acids within the first α -helix;
 - (iii) one or more exposed amino acids in said α-helix which have acidic or acidic-like properties being substituted by a basic amino acid residue;
 - (iv) being in recombinant or synthetic form;
 - (v) being capable of acting as an antagonist for at least one property of the corresponding native HGF.
- This aspect of the present invention is predicated in part on the surprising discovery that a mutation in amino acid 21 (Glu) of hGM-CSF to Arg or Lys together with the variant in GM-CSF being in unglycosylated form results in the hGM-CSF variant being unable to detectably exhibit classical GM-CSF function. The variants, referred to herein "GM-CSF Arg²¹" and "GM-CSF Lys²¹" are unable to bind to high affinity receptors but are still able to fully bind the low affinity α chain of the GM-CSF receptor. Importantly, the non-glycosylated GM-CSF Arg²¹ and GM-CSF Lys²¹ act as antagonists, preventing the stimulatory effect of native GM-CSF. For convenience, the numbering of amino acid

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residues in hGM-CSF is taken from Wong et al. (1985).

By way of a shorthand notation the following three letter abbreviations for amino acid residues are used in the specification as defined in Table 1.

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Where a specific amino residue is referred to by its position in the polypeptide of an HGF, the amino acid abbreviation is used with the residue number given in superscript (i.e. Xaaⁿ, wherein Xaa is the amino acid residue).

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Table 1

15	Amino acid	Three-letter abbreviation	Corresponding single-letter abbreviation
	Alanine	Ala	Α
	Arginine	Arg	R
20	Asparagine	Asn	N
	Aspartic acid	Asp	D
	Cysteine	Cys	C
	Glutamine	Gln	Q
	Glutamic acid	Glu	E
25	Glycine	Gly	G
	Histidine	His	Н
	Isoleucine	Ile	I
	Leucine	Leu	L
	Lysine	Lys	K
30	Methionine	Met	M
	Phenylalanine	Phe ·	F
	Proline	Pro	P
	Serine	Ser	S
	Threonine	Thr	T
35	Tryptophan	Trp	W
	Tyrosine	Tyr	Y
	Valine	Val	V

The present invention is exemplified using GM-CSF and in particular hGM-CSF Arg²¹ and hGM-CSF Lys²¹. This is done, however, with the understanding that the present invention extends to all HGFs as hereinbefore described.

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For example, given that there is a Glu at position 22 of IL-3 and position 13 in IL-5 which in the three dimensional structure (in the case of IL-5) occupies an equivalent position to Glu^{21} of GM-CSF, then the present invention provides the basis for the creation of antagonists in IL-3 and IL-5. The substitution of Glu at these positions in IL-3 and IL-5 may be the sole mutation or it may be in combination with other amino acid mutations (including substitutions, deletions and/or additions) for the development of effective antagonists. This similarly applies to other HGFs based on the acidic or acidic-like amino acid residues in the first α -helix of the molecule. The location of the N-terminal helix can in each case be readily determined on comparable motifs from predicted helices. Such HGFs are listed in Table 2 showing the acidic or acidic-like amino acid residue in bold in the equivalent position to Glu^{21} of hGM-CSF. The acidic or acidic-like amino acid residues are readily substituted by, for example, recombinant DNA technology.

- According to another aspect of the present invention there is provided a modified variant including HGF comprising an amino acid sequence in the first α-helix of said HGF selected from the group consisting of:
- i) His Val Asn Ala Ile Gln Xaa Ala Arg Arg Leu Leu Asn Leu (SEQ ID No. 1);
 - ii) Ala Leu Val Lys Xaa Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu (SEQ ID No. 2);
 - iii) Asn Met Ile Xaa Xaa Ile Ile Thr His Leu (SEQ ID No. 3);
 - iv) Leu Leu Xaa Leu Gln Met Ile Leu (SEQ ID No. 4);
- 30 v) Ile Thr Leu Gln Xaa Ile Ile Lys Thr Leu (SEQ ID No. 5);
 - vi) Arg Tyr Ile Leu Xaa Gly Ile Ser Ala Leu Arg Lys
 (SEQ ID No. 6);
 - vii) Gly Asp Gln Tyr Xaa Ser Val Leu Met Val Ser Ile
 (SEQ ID No. 7);

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- viii) Ala Gly Ile Leu Xaa Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp (SEQ ID No. 8);
- ix) Asn Met Leu Arg Xaa Leu Arg Asp Ala Phe Ser (SEQ ID No. 9);
- 5 x) Phe Leu Leu Lys Cys Leu Xaa Gln Val Arg Lys Ile (SEQ ID No. 10); and
 - xi) Tyr Leu Leu Glu Ala Lys Xaa Ala Glu Asn Ile Thr Thr Gly (SEQ ID No. 11);
- wherein Xaa is a basic amino acid, preferably selected from the group consisting of Arg and Lys, and wherein said variant HGF is in unglycosylated form and acts as an antagonist for at least one property of the corresponding native HGF. Preferably, the haemopoietic growth factor is hGM-CSF and Xaa is Arg or Lys at position 21 of the first α-helix.

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- The HGF antagonists of the present invention and in particular GM-CSF Arg²¹ and GM-CSF Lys²¹ are useful *inter alia* in the treatment of myeloid and lymphocyte leukaemias, some tumours of non-haemopoietic origins and acute and chronic inflammation such as asthma and rheumatoid arthritis. These and other conditions are considered herein to result from or be facilitated by the aberrant effects of an endogenous HGF such as GM-CSF. hGM-CSF Arg²¹ and hGM-CSF Lys²¹ will also be useful in mobilising stem cells and progenitor cells into the circulation without the risk of activating neutrophils and monocytes. Other related molecules may have different useful properties.
- The present invention, therefore, contemplates a method of treatment comprising the administration to a mammal of an effective amount of a modified HGF as hereinbefore defined and in particular hGM-CSF Arg²¹ or hGM-CSF Lys²¹ or both for a time and under conditions sufficient for effecting said treatment. Generally, the mammal is a human, livestock animal, companion animal or laboratory test animal. Most preferably, the mammal is a human. A single modified HGF may be administered or a combination of variants of the same HGF. For example, a range of hGM-CSF variants could be used such as a combination of hGM-CSF Arg²¹ and hGM-CSF Lys²¹.

TABLE 2

Cytokines related to GM-CSF exhibit a conserved acidic residue analogous to E21 in GM-CSF

CYTOKINE ²	HEI (Amir Resid	HELIX ¹ (Amino Acid Residue No.)						EM .	7 ON:	CID	SEQU	AMINO ACID SEQUENCE	6						SEQ ID No.
hGM-CSF	15-28	His Val	Asn	Ala	Ile	Gln	Glu	Ala	Glu Ala Arg	Arg	Leu	Leu Leu Asn	Asn	Leu					12
hIL-5	9-24		Ala	Leu	val	Lys	Glu	Thr	Leu	Ala	Leu	Leu Leu	Ser	Thr	His	His Arg Thr	Thr 1	Leu	13
hIL-3	18-27		Asn	Met	: Ile	ABP	Gla	Ile	Ile	Thr	His	Leu							14
hIL-2	17-25			Leu	Leu	Leu	Asp	Leu	Gln	Met	Ile	Leu							15
hIL-4	8-17		Ile	Thr	. Leu	Gln	Asp	Ile	11e	Lys	$_{\mathrm{Thr}}$	Leu							16
hIL-6	18-43		Arg	Tyr	· Ile	Leu	Asp	Gly	Ile	Ser	Ala	Leu Arg		Lys					17
hIL-7	9-20		Gly	, Asp	Glu	Tyr	Gla	Ser	Val	Leu	Met	Val	Ser	Ile					18
hIL-9	7-22	•	Ala	Gly	, Ile	Leu	ABP	Ile	Asn	Phe	Leu	Ile	Asn	Lув	Met	Gln	glu i	Asp	19
hIL-10	21-31		Asn	Met	ren:	Arg	Asp	Leu	Arg	Asp	Ala	Phe	Ser						50
hG-CSF	13-24	Phe Leu	r Leu	LyB	Cys	Leu	Gla	Gln	Val	Arg	Lys	Ile							21
hEPO	4-28	Tvr Leu	Len	Glu	Ala	Lys	Glu	Ala	Glu	Asn	Ile	Thr	Thr	Gly	-				22

Only the pertinent portion of each helix is shown.

The location of the N-terminal helix in the Zurawski and Zurawski, 1989); predicted helices are from: for IL-2, (Brandhuber et al, 1987; Zurawski and Zurawski, 1985 for hIL-3 (Parry et al, 1988); for mIL-5, (Parry et al, 1988); for hIL-6 (Bazan, 1990a); for hG-CSF, (Parry et al, 1988); for hBPO, (Bazan, 1990); for hGM-CSF the first helix was other cytokines was based on comparable motifs from these secondary structure predictions. determined from the crystal structure (Karplus, 1991).

The present invention also provides a pharmaceutical composition comprising the variant HGFs as hereinbefore defined or combinations thereof. Most particularly, the pharmaceutical composition comprises hGM-CSF Arg²¹ or hGM-CSF Lys²¹ or both.

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Methods for preparing pharmaceutical compositions are well known in the art and reference can conveniently be made to Remington's Pharmaceutical Sciences, Mack Publishing Company, Eaton, Pennsylvania, USA and may also include one or more pharmaceutical acceptable carriers and/or diluents.

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The present invention is further described by reference to the following non-limiting Examples and/or Figures.

In the Figures:

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Figure 1 is a graphical representation showing titration of E. coli derived GM-CSF Arg^{21} for its ability to affect 0_2^- production in human neutrophils (\square) and to antagonise the enhancement of 0_2^- by wild type GM-CSF tested at 1.0 ng/ml (\triangle) and at 0.1 ng/ml (closed triangle).

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- Figure 2 is a graphical representation showing failure of E. coli-derived GM-CSF Arg^{21} to antagonise the enhancement of 0_2 production in human neutrophils stimulated with tumour necrosis factor- α (TNF- α).
- Figure 3 is a graphical representation showing competitive inhibition of 125 I-GM-CSF binding to COS cells transfected with the GM-CSF receptor α chain alone (top) or α and β chains (bottom) by GM-CSF Arg²¹.
- Figure 4 is a graphical representation showing titration of GM-CSF Arg²¹ for its ability to antagonise GM-CSF (A) in contrast to no effect on IL-3 (B)-mediated proliferation of TF-1 cells.

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Figure 5 is a graphical representation showing the titration of GM-CSF Arg²¹ for its ability to antagonise TF-1 proliferation stimulated by either *E. coli*-derived GM-CSF (A), yeast-derived GM-CSF (B) or CHO-derived GM-CSF (C).

Figure 6 is a graphical representation showing the titration of GM-CSF Arg²¹ for its ability to antagonise three primary human myeloid leukaemia (A, B & C) ex vivo.

Figure 7 is a graphical representation showing that E21R antagonises both GM-CSF but not TNF- α -mediated stimulation of human neutrophils (A), and both E21R and E21K also antagonise neutrophil stimulation by CHO cell-derived GM-CSF (B). In panel A, titrations of E. coli-derived wild type GM-CSF (\bullet), TNF- α (\bullet) and E21R (\blacksquare) are shown. In antagonistic experiments, E21R was titrated against 1ng/ml of E. coli-derived GM-CSF (\square) or 3ng/ml TNF- α (\diamond). In panel B, titrations of CHO cell-derived wild type GM-CSF (\square) and E21R (\blacksquare) and E21K (\triangle) are shown. In antagonistic experiments, E21R (\square) or E21K (\triangle) were titrated against 3ng/ml CHO cell-derived GM-CSF. Each value represents the mean of triplicate determinations and error bars represent the SEM.

EXAMPLE 1

Expression of wild type and GM-CSF Arg²¹ and GM-CSF Lys²¹ in an *E. coli* expressed system

Wild type GM-CSF was expressed in *E. coli* using a plasmid (designated pshGM-CSF) containing a synthetic human GM-CSF cDNA cloned into the *E. coli* expression vector pIN-III-OmpH3, a derivative of the vector pIN-III-OmpA2 (Ghrayeb *et al*, 1984). GME21R was expressed from the plasmid pSGM21.1 containing $Glu^{21} \rightarrow Arg^{21}$ substitution and was derived from the pSGM-CSF parental plasmid. GME21K was expressed from the plasmid pSGM21.4 containing $Glu^{21} \rightarrow Lys^{21}$ substitution and was derived from the pSGM-CSF parental plasmid.

pSGM21 was generated by initially eliminating a SacII site from the wild type GM-CSF using oligonucleotide cassette mutagenesis to generate plasmid pSGMV1. A 64 bp Nco 1/SacII fragment was then excised from the pSGMV1 plasmid and replaced by double-stranded 64bp oligonucleotides containing the appropriate mutation in the DNA sequence.

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pSGM21.4 was generated by excising an 88 bp Bgl11/SacII fragment from pSGF1 and replacing it with 88bp oligonucleotides containing the appropriate mutation site directed mutagenesis (Zoller & Smith, 1984).

5 Protein was expressed in either MC1061, for wild type GM-CSF or BL21 for GME21R or GME21K, after induction by isopropyl β-D-thiogalactoside and recovered from the periplasmic space by osmotic shock (Koshland and Botstein, 1980).

GM-CSF protein was purified using a monoclonal antibody 4A12 generated in the laboratory coupled to Sepharose beads. Further purification was achieved by reversed phase HPLC using a BioRad controller and a Brownlee Aquaport C8 100 x 10mm column. GM-CSF was eluted using a 30-50% gradient of acetonitrile in 0.1% trifluoroacetic acid.

Resultant purified GM-CSF was lyophilised and resuspended in 1 x PBS before being quantitated by HPLC gel filtration. Samples were fractionated on a Beckman Ultraspherogel SEC3000 7.5 x 300mm using a 0.1M Na Phosphate pH 7.0/01M Na₂SO₄ mobile phase. Purity was estimated at >95% and area under peaks corresponding to GM-CSF integrated as the extinction coefficient of 0.95 absorbance units.ml.mg⁻¹.

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EXAMPLE 2

Visualisation of mutant GM-CSF protein

GM-CSF unpurified or purified from *E. coli* was size-fractionated by NaDodSo₄/12.5% w/v polycarylamide gel electrophoresis (Laemmli, 1970). For Western blot analysis, protein was transferred to nitrocellulose as described (Towbin *et al*, 1979). Filters were probed with a sheep anti-GM-CSF followed by a second layer of biotinylated-rabbit anti-sheep IgG. After a further incubation with an avidin-biotinylated-horseradish peroxidase conjugate, the complex was visualised using a diaminobenzidine substrate solution. For silver staining, the method of Morrissey (1981) was used.

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EXAMPLE 3

Stimulation of haemopoietic cell proliferation

The human erythroleukaemia cell line TF-1 (and myeloid leukaemia) cells were used to measure the proliferative function of GM-CSF and GM-CSF Arg²¹. Proliferation of TF-1 cells were measured by the ability to incorporate [³H]-thymidine in response to increasing doses of GM-CSF. This assay was performed as described by Lopez *et al* (1988).

EXAMPLE 4

Functional activation of human granulocytes and monocytes

The superoxide anion production assay was carried out as previously described (Lopez et al, 1986).

EXAMPLE 5

(a) Radioiodination of GM-CSF.

Radioreceptor assay

Yeast derived human GM-CSF or *E. coli*-derived human GM-CSF was radioiodinated by the ICl method (Contreras *et al*, 1983). Iodinated protein was separated from free ¹²⁵I by chromatography on a Sephadex G-25 PD10 column (Pharmacia, Uppsala, Sweden), equilibrated in phosphate buffered saline (PBS) containing 0.02% w/v Tween 20, and stored at 4°C for up to 4 weeks. Before use, the iodinated protein was purified from Tween and non-protein-associated radioactivity by cation exchange chromatography on a 0.3ml CM-Sepharose CL-6B column (Pharmacia) and stored at 4°C for up to 5 days. The radiolabelled GM-CSF retained >90% biological activity as judged from titration curves using non-iodinated GM-CSF as controls.

(b) Competition binding assays.

Competition for binding to high affinity and low affinity receptors used stably transfected CHO cell lines expressing either the α and β chains, or the α chain alone. The cells were suspended in binding medium consisting of RPMI-1640 supplemented with 20mmol/l HEPES and 0.5% w/v bovine serum albumin (BSA) and 0.1% w/v sodium azide. Typically, equal volumes (50µl) of 4 x 10⁴ CHO cells, iodinated GM-CSF and different

concentrations of GM-CSF and GM-CSF Arg²¹ were mixed in siliconised glass tubes for 3 hr at 4°C. At the end of the incubation period, cell suspensions were overlaid on 0.2ml foetal calf serum (FCS) at 4°C, centrifuged in a Beckman Microfuge 12, and the tip of each tube containing the visible cell pellet cut off and counted in a gamma counter. Specific counts were determined by first subtracting the counts, obtained in the presence of excess wild type GM-CSF.

EXAMPLE 6

Generation of hGM-CSF Variants

- By way of example only, the generation of GM-CSF Arg²¹ is hereinafter described in detail. A human GM-CSF cDNA was subjected to mutagenesis to introduce the amino acid Arg for Glu at position 21. Two mutants were obtained, one containing the Glu²¹→Arg mutation and a second one containing a double mutation X¹⁰→Ile and Glu²¹→Arg. These mutants were cloned into the expression system piN OMPIII and expressed in E. coli. Wild type (WT) GM-CSF was expressed in MC1061. GM-CSF Arg²¹ could not be expressed in MC1061. Of twenty strains tested for GM-CSF Arg²¹ expression, BL21 was the highest producer and used for subsequent studies.
 - To obtain purified GM-CSF Arg²¹ in high yields a two-step purification procedure was devised. In the first step, GM-CSF Arg²¹ was purified by an affinity column constructed with a monoclonal antibody (4A12) that binds to GM-CSF in solution and with high affinity. Affinity-purified GM-CSF Arg²¹ was then purified by reverse-phase HPLC and quantitated by HPLC before being analysed for biological and binding activities.
- It was found that *E. coli*-derived GM-CSF Arg²¹ was unable to enhance neutrophil 0₂production up to a concentration of 3,000 ng/ml (Figure 1). This is different to the
 inventors' previous results with CHO-derived (i.e. glycosylated) GM-CSF Arg²¹ which
 was able to enhance neutrophil function at approximately 30 ng/ml which represents a
 300 fold reduced potency compared to wild type GM-CSF (Lopez *et al*, 1992).

Despite its inability to activate neutrophils, *E. coli*-derived GM-CSF Arg^{21} was able to bind as well as wild type GM-CSF to the α -chain of the GM-CSF receptor (Figure 3). In contrast, GM-CSF Arg^{21} exhibited an approximate 100-fold reduction in binding to the $\alpha\beta$ GM-CSF receptor complex (Figure 3) indicating that the influence of the β chain has been selectively lost.

This is the first time that a GM-CSF mutant is shown to have unaltered binding to the GM-CSF receptor α -chain as well as being devoid of agonistic activity. This indicates that whilst binding to the α chain is necessary for GM-CSF activity it is not sufficient, and that GM-CSF binding to the β chain is required for GM-CSF-mediated activation.

Since $E.\ coli$ -derived GM-CSF Arg^{21} was not able to stimulate neutrophils yet fully bound the GM-CSF receptor α chain, it was tested for antagonistic activity. The inventors found that this mutant fully inhibited the effect of wild type GM-CSF with an approximately 300-fold excess required to induce 50% inhibition of $E.\ coli$ -derived wild type GM-CSF (Figure 1). This antagonistic effect was specific for GM-CSF as judged by the lack of antagonistic effect of GM-CSF Arg^{21} on TNF enhancement of neutrophil 0_2 - production (Figure 2).

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The antagonistic effect of *E. coli*-derived GM-CSF Arg²¹ was present whether wild type GM-CSF was expressed in *E. coli* (Figure 5A), yeast (Figure 5B) or CHO cells (Figure 5C). In keeping with the differences in binding affinity between heavily glycosylated CHO GM-CSF (lower affinity), partially glycosylated yeast GM-CSF (intermediate affinity) and unglycosylated *E. coli* GM-CSF (higher affinity), *E. coli*-derived GM-CSF Arg²¹ antagonised better the CHO wild type GM-CSF (Figure 5).

The antagonism of *E. coli*-derived GM-CSF Arg²¹ was not restricted to proliferation of the established TF-1 cell line but was also seen in primary myeloid leukaemias. In three different leukaemias, *E. coli*-derived GM-CSF Arg²¹ antagonised the proliferative effect of wild type *E. coli* GM-CSF with an EC50 that varied with each leukaemia (Figure 6).

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These results show that an unglycosylated GM-CSF molecule with a mutated Glu for an Arg in position 21 of the first α -helix is able to antagonise native GM-CSF.

Since the mutated Glu in GM-CSF is in a position where the same acidic residue or similar acidic residues (e.g. Asp) are present in related growth factors (see Table 2), this invention extends to antagonistic molecules for these growth factors constructed by incorporating the analogous charge reversal mutation. In particular, given that the GM-CSF, IL-3 and IL-5 receptor share the β chain, the Glu²² in IL-3 and the Glu¹³ in IL-5 are predicted to play a similar role. Other variant HGFs are shown in Examples 3 to 22 in which the equivalent or similar amino acid residue to Glu²¹ of hGM-CSF is replaced by either Arg or Lys. In these Examples, the amino acid sequences are provided for the relevant portion of the first α -helix carrying the substitution (see Table 2).

EXAMPLE 7 hGM-CSF Lvs²¹

nGM-CSF Lys

His Val Asn Ala Ile Gln Lys Ala Arg Arg Leu Leu Asn Leu (SEQ ID NO. 23)

EXAMPLE 8

hlL-5 Lys¹³

20 Ala Leu Val Lys Lys Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu (SEQ ID NO.24)

EXAMPLE 9

hlL-5 Arg¹³

Ala Leu Val Lys Arg Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu (SEQ ID NO. 25)

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EXAMPLE 10

hIL-3 Variants

	Asn Met Ile Asp Lys Ile Ile Thr His Leu	hIL-3 Lys ²² (SEQ ID NO. 26)
30	Asn Met Ile Lys Glu Ile Ile Thr His Leu	hIL-3 Lys ²¹ (SEQ ID NO. 27)
	Asn Met Ile Asp Arg Ile Ile Thr His Leu	hIL-3 Arg ²² (SEQ ID NO. 28)
	Asn Met Ile Arg Glu Ile Ile Thr His Leu	hIL-3 Arg ²¹ (SEQ ID NO. 29)
	Asn Met Ile Lys Lys Ile Ile Thr His Leu	hIL-3 Lys ²¹ Lys ²² (SEQ ID NO. 30)

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Asn Met Ile Arg Arg Ile Ile Thr His Leu

hIL-3 Arg²¹ Arg²² (SEQ ID NO. 31)

EXAMPLE 11

hIL-2 Lys²⁰

5 Leu Leu Leu Lys Leu Gln Met Ile Leu (SEQ ID NO. 32)

EXAMPLE 12

hIL-2 Arg²⁰

Leu Leu Leu Arg Leu Gln Met Ile Leu (SEQ ID NO. 33)

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EXAMPLE 13

hIL-4 Lys¹²

Ile Thr Leu Gln Lys Ile Ile Lys Thr Leu (SEQ ID NO. 34)

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EXAMPLE 14

hIL-4 Arg¹²

Ile Thr Leu Gln Arg Ile Ile Lys Thr Leu (SEQ ID NO. 35)

EXAMPLE 15

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hIL-6 Lys²²

Arg Tyr Ile Leu Lys Gly Ile Ser Ala Leu Arg Lys (SEQ ID NO. 36)

EXAMPLE 16

hIL-6 Arg²²

25 Arg Tyr Ile Leu Arg Gly Ile Ser Ala Leu Arg Lys (SEQ ID NO. 37)

EXAMPLE 17

hIL-7 Lys¹³

Gly Asp Gln Tyr Lys Ser Val Leu Met Val Ser Ile (SEQ ID NO. 38)

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EXAMPLE 18

hIL-7 Arg¹³

Gly Asp Gln Tyr Arg Ser Val Leu Met Val Ser Ile (SEQ ID NO. 39)

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EXAMPLE 19

hIL-9 Lys¹¹

Ala Gly Ile Leu Lys Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp (SEQ ID NO. 40)

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EXAMPLE 20

hIL-9 Arg¹¹

Ala Gly Ile Leu Arg Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp (SEQ ID NO. 41)

EXAMPLE 21

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hIL-10 Lys²⁵

Asn Met Leu Arg Lys Leu Arg Asp Ala Phe Ser (SEQ ID NO. 42)

EXAMPLE 22

hIL-10 Arg²⁵

20 Asn Met Leu Arg Arg Leu Arg Asp Ala Phe Ser (SEQ ID NO. 43)

EXAMPLE 23

hG-CSF Lys¹⁹

Phe Leu Leu Lys Cys Leu Lys Gln Val Arg Lys Ile (SEQ ID NO. 44)

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EXAMPLE 24

hG-CSF Arg¹⁹

Phe Leu Leu Lys Cys Leu Arg Gin Val Arg Lys Ile (SEQ ID NO. 45)

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EXAMPLE 25 hEPO Lys¹⁰

Tyr Leu Leu Glu Ala Lys Lys Ala Glu Asn Ile Thr Thr Gly (SEQ ID NO. 46)

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EXAMPLE 26 hEPO Arg¹⁰

Tyr Leu Leu Glu Ala Lys Arg Ala Glu Asn Ile Thr Thr Gly (SEQ ID NO. 47)

Those skilled in the art will appreciate that the invention described herein is susceptible to variations and modifications other than those specifically described. It is to be understood that the invention includes all such variations and modifications. The invention also includes all of the steps, features, compositions and compounds referred to or indicated in this specification, individually or collectively, and any and all combinations of any two or more of said steps or features.

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SEQUENCE LISTING

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 - (B) COMPUTER: IBM PC compatible
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 - (D) SOFTWARE: Patentin Release #1.0, Version #1.25
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- (2) INFORMATION FOR SEQ ID NO. 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14 amino acid residues
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 1:

His Val Asn Ala Ile Gln Xaa Ala Arg Arg Leu Leu Asn Leu

- (2) INFORMATION FOR SEQ ID NO. 2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 2:

Ala Leu Val Lys Xaa Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10 amino acid residues
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 3:

Asn Met Ile Xaa Xaa Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 4:

Leu Leu Xaa Leu Gln Met Ile Leu

- (2) INFORMATION FOR SEQ ID NO. 5:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 5:

Ile Thr Leu Gln Xaa Ile Ile Lys Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 6:

Arg Tyr Ile Leu Xaa Gly Ile Ser Ala Leu Arg Lys

- (2) INFORMATION FOR SEQ ID NO. 7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 7:

Gly Asp Gln Tyr Xaa Ser Val Leu Met Val Ser Ile

- (2) INFORMATION FOR SEQ ID NO. 8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 8:

Ala Gly Ile Leu Xaa Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp

- (2) INFORMATION FOR SEQ ID NO. 9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 9:

Asn Met Leu Arg Xaa Leu Arg Asp Ala Phe Ser

- (2) INFORMATION FOR SEQ ID NO. 10:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 10:

Phe Leu Leu Lys Cys Leu Xaa Gln Val Arg Lys Ile

- (2) INFORMATION FOR SEQ ID NO. 11:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 11:

Tyr Leu Leu Glu Ala Lys Xaa Ala Glu Asn Ile Thr Thr Gly

- (2) INFORMATION FOR SEQ ID NO. 12:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 12:

His Val Asn Ala Ile Gln Glu Ala Arg Arg Leu Leu Asn Leu

- (2) INFORMATION FOR SEQ ID NO. 13:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 13:

Ala Leu Val Lys Glu Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 14:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 14:

Asn Met Ile Asp Glu Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 15:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 15:

Leu Leu Leu Asp Leu Gln Met Ile Leu

- INFORMATION FOR SEQ ID NO. 16: (2)
 - SEQUENCE CHARACTERISTICS: (i)
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - STRANDEDNESS: single (C)
 - (D) TOPOLOGY: linear

 - (ii) MOLECULE TYPE: polypeptide
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 16:

Ile Thr Leu Gln Asp Ile Ile Lys Thr Leu

- INFORMATION FOR SEQ ID NO. 17: (2)
 - (i) SEQUENCE CHARACTERISTICS:
 - LENGTH: 12 (A)
 - TYPE: amino acid (B)
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 17:

Arg Tyr Ile Leu Asp Gly Ile Ser Ala Leu Arg Lys

- INFORMATION FOR SEQ ID NO. 18: (2)
 - SEQUENCE CHARACTERISTICS: (i)
 - (A) LENGTH: 12
 - TYPE: amino acid (B)
 - (C) STRANDEDNESS: single
 - TOPOLOGY: linear (D)
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 18:

Gly Asp Gln Tyr Glu Ser Val Leu Met Val Ser Ile

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- (2) INFORMATION FOR SEQ ID NO. 19:
 - SEQUENCE CHARACTERISTICS: (i)
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - STRANDEDNESS: single (C)
 - (D) TOPOLOGY: linear
 - MOLECULE TYPE: polypeptide (ii)
 - SEQUENCE DESCRIPTION: SEQ ID NO. 19:

Ala Gly Ile Leu Asp Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp

- (2) INFORMATION FOR SEQ ID NO. 20:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 20:

Asn Met Leu Arg Asp Leu Arg Asp Ala Phe Ser

- (2) INFORMATION FOR SEQ ID NO. 21:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 21:

Phe Leu Leu Lys Cys Leu Glu Gln Val Arg Lys Ile

- (2) INFORMATION FOR SEQ ID NO. 22:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 22:

Tyr Leu Leu Glu Ala Lys Glu Ala Glu Asn Ile Thr Thr Gly

- (2) INFORMATION FOR SEQ ID NO. 23:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 23:

His Val Asn Ala Ile Gln Lys Ala Arg Arg Leu Leu Asn Leu

- (2) INFORMATION FOR SEQ ID NO. 24:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 24:

Ala Leu Val Lys Lys Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 25:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 25:

Ala Leu Val Lys Arg Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 26:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 26:

Asn Met Ile Asp Lys Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 27:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 27:

Asn Met Ile Lys Glu Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 28:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 28:

Asn Met Ile Asp Arg Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 29:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 29:

Asn Met Ile Arg Glu Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 30:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 30:

Asn Met Ile Lys Lys Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 31:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 31:

Asn Met Ile Arg Arg Ile Ile Thr His Leu

- (2) INFORMATION FOR SEQ ID NO. 32:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 32:

Leu Leu Lys Leu Gln Met Ile Leu

- (2) INFORMATION FOR SEQ ID NO. 33:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 33:

Leu Leu Arg Leu Gln Met Ile Leu

- (2) INFORMATION FOR SEQ ID NO. 34:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 34:

Ile Thr Leu Gln Lys Ile Ile Lys Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 35:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 35:

Ile Thr Leu Gln Arg Ile Ile Lys Thr Leu

- (2) INFORMATION FOR SEQ ID NO. 36:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 36:

Arg Tyr Ile Leu Lys Gly Ile Ser Ala Leu Arg Lys

- (2) INFORMATION FOR SEQ ID NO. 37:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 37:

Arg Tyr Ile Leu Arg Gly Ile Ser Ala Leu Arg Lys

- (2) INFORMATION FOR SEQ ID NO. 38:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 38:

Gly Asp Gln Tyr Lys Ser Val Leu Met Val Ser Ile

- (2) INFORMATION FOR SEQ ID NO. 39:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 39:

Gly Asp Gln Tyr Arg Ser Val Leu Met Val Ser Ile

- (2) INFORMATION FOR SEQ ID NO. 40:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 40:

Ala Gly Ile Leu Lys Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp

- (2) INFORMATION FOR SEQ ID NO. 41:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 16
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 41:

Ala Gly Ile Leu Arg Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp

- (2) INFORMATION FOR SEQ ID NO. 42:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 42:

Asn Met Leu Arg Lys Leu Arg Asp Ala Phe Ser

- (2) INFORMATION FOR SEQ ID NO. 43:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 43:

Asn Met Leu Arg Arg Leu Arg Asp Ala Phe Ser

- (2) INFORMATION FOR SEQ ID NO. 44:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 44:

Phe Leu Leu Lys Cys Leu Lys Gln Val Arg Lys Ile

- (2) INFORMATION FOR SEQ ID NO. 45:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 45:

Phe Leu Leu Lys Cys Leu Arg Gln Val Arg Lys Ile

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- (2) INFORMATION FOR SEQ ID NO. 46:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 46:

Tyr Leu Leu Glu Ala Lys Lys Ala Glu Asn Ile Thr Thr Gly

- (2) INFORMATION FOR SEQ ID. NO. 47:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 14
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: polypeptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO. 47:

Tyr Leu Leu Glu Ala Lys Arg Ala Glu Asn Ile Thr Thr Gly

WO 95/04075 PCT/AU94/00432

CLAIMS:

1. A modified haemopoietic growth factor (HGF) characterised by being in unglycosylated form and comprising a sequence of amino acids within a first \alpha-helix wherein one or more exposed amino acids in said first α-helix having acidic or acidic-like properties are substituted with a basic amino acid residue.

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- 2. A modified HGF according to claim 1, wherein said HGF is a modified form of an HGF selected from granulocyte-macrophage colony-stimulating factor (GM-CSF), interleukin (IL) -2, IL-3, IL-4, IL-5, IL-6, Il-7, IL-9, IL-10, G-CSF, erythropoietin (EPO).
- A modified HGF according to claim 2 wherein said HGF is a modified form of 3. GM-CSF.
- A modified HGF according to claim 1 or 2 or 3 wherein said HGF is of human, livestock animal, companion animal or laboratory test animal origin.
- 5. A modified HGF according to claim 4 wherein said HGF is of human origin.
- 6. A modified HGF according to any one of the preceding claims wherein the acidic amino acid residue on the first α-helix is Glu and/or Asp and the basic amino acid residue substituted therefor is Arg and/or Lys.
- 7. A modified HGF comprising an amino acid sequence in the first α-helix of said HGF selected from the group consisting of:
- i) His Val Asn Ala Ile Gln Xaa Ala Arg Arg Leu Leu Asn Leu;
- ii) Ala Leu Val Lys Xaa Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu;
- iii) Asn Met Ile Xaa Xaa Ile Ile Thr His Leu;
- iv) Leu Leu Leu Xaa Leu Gln Met Ile Leu;
- v) Ile Thr Leu Gln Xaa Ile Ile Lys Thr Leu;

- vi) Arq Tyr Ile Leu Xaa Gly Ile Ser Ala Leu Arg Lys;
- vii) Gly Asp Gln Tyr Xaa Ser Val Leu Met Val Ser Ile;
- viii) Ala Gly Ile Leu Xaa Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp;
- ix) Asn Met Leu Arg Xaa Leu Arg Asp Ala Phe Ser;
- x) Phe Leu Leu Lys Cys Leu Xaa Gln Val Arg Lys Ile;
- xi) Tyr Leu Leu Glu Ala Lys Xaa Ala Glu Asn Ile Thr Thr Gly;

wherein Xaa is a basic amino acid, selected from the group consisting of Arg and Lys and wherein said modified HGF is in unglycosylated form and acts as an antagonist for at least one property of the corresponding native HGF.

- 8. A modified HGF according to claim 7 wherein the HGF is a modified form of an HGF selected from the list consisting of GM-CSF, IL-2, IL-3, IL-4, IL-5, IL-6, Il-7, IL-9, IL-10, G-CSF, EPO.
- 9. A modified HGF according to claim 8, wherein the variant HGF is a modified form of GM-CSF.
- 10. A modified HGF according to claim 7 or 8 wherein Xaa is Arg.
- 11. A modified human GM-CSF in unglycosylated form having an amino acid sequence in a first α -helix comprising His Val Asn Ala Ile Gln Arg Ala Arg Arg Leu Leu Asn Leu.
- 12. A modified human GM-CSF in unglycosylated form having an amino acid sequence in a first α -helix comprising His Val Asn Ala Ile Gln Lys Ala Arg Arg Leu Leu Asn Leu.
- 13. A modified human IL-5 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Ala Leu Val Lys Lys Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu.

14. A modified human IL-5 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Ala Leu Val Lys Arg Thr Leu Ala Leu Leu Ser Thr His Arg Thr Leu.

15. A modified human IL-3 in unglycosylated form having an amino acid sequence in a first α -helix selected from the list consisting of:

Asn Met Ile Asp Lys Ile Ile Thr His Leu;

Asn Met Ile Lys Glu Ile Ile Thr His Leu;

Asn Met Ile Asp Arg Ile Ile Thr His Leu;

Asn Met Ile Arg Glu Ile Ile Thr His Leu;

Asn Met Ile Lys Lys Ile Ile Thr His Leu; and

Asn Met Ile Arg Arg Ile Ile Thr His Leu.

16. A modified human IL-2 in unglycosylated form having an amino acid sequence in a first α-helix comprising

Leu Leu Lys Leu Gln Met Ile Leu.

17. A modified human IL-2 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Leu Leu Leu Arg Leu Gln Met Ile Leu.

18. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Ile Thr Leu Gln Lys Ile Ile Lys Thr Leu.

19. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Ile Thr Leu Gln Arg Ile Ile Lys Thr Leu.

20. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Arg Tyr Ile Leu Lys Gly Ile Ser Ala Leu Arg Lys.

21. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Arg Tyr Ile Leu Arg Gly Ile Ser Ala Leu Arg Lys.

22. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Gly Asp Gln Tyr Lys Ser Val Leu Met Val Ser Ile.

23. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Gly Asp Gln Tyr Arg Ser Val Leu Met Val Ser Ile.

24. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α-helix comprising

Ala Gly Ile Leu Lys Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp.

25. A modified human IL-4 having an amino acid sequence in a first α -helix comprising

Ala Gly Ile Leu Arg Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp.

26. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Asn Met Leu Arg Lys Leu Arg Asp Ala Phe Ser.

27. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α-helix comprising

Asn Met Leu Arg Arg Leu Arg Asp Ala Phe Ser.

28. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Phe Leu Leu Lys Cys Leu Lys Gln Val Arg Lys Ile.

29. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Phe Leu Lys Cys Leu Arg Gln Val Arg Lys Ile.

30. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

Tyr Leu Leu Glu Ala Lys Lys Ala Glu Asn Ile Thr Thr Gly.

31. A modified human IL-4 in unglycosylated form having an amino acid sequence in a first α -helix comprising

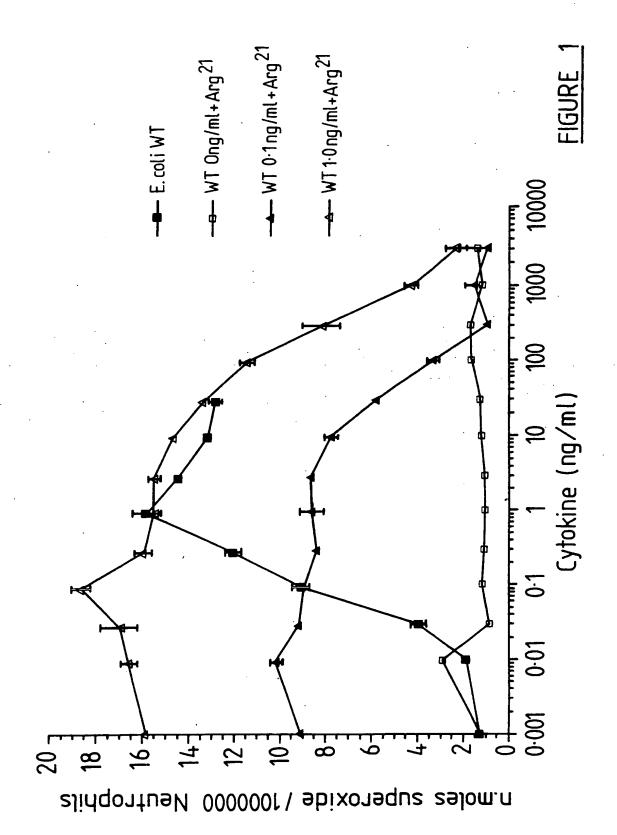
Tyr Leu Leu Glu Ala Lys Arg Ala Glu Asn Ile Thr Thr Gly.

- 32. A method of ameliorating the aberrant effects of an endogenous HGF in a mammal, said method comprising administering to said mammal an effective amount of a modified HGF characterised by being in unglycosylated form and comprising a sequence of amino acids within a first α -heliz wherein one or more exposed amino acids in said first α -helix having acidic or acidic-like properties are substituted with a basic amino acid residue.
- 33. A method according to claim 32 wherein the modified HGF is selected from the group consisting of:
- i) His Val Asn Ala Ile Gln Xaa Ala Arg Arg Leu Leu Asn Leu;
- ii) Ala Leu Val Lys Xaa Thr Leu Ala Leu Leu Ser Thr His Arg
 Thr Leu;
- iii) Asn Met Ile Xaa Xaa Ile Ile Thr His Leu;
- iv) Leu Leu Leu Xaa Leu Gln Met Ile Leu;
- v) Ile Thr Leu Gln Xaa Ile Ile Lys Thr Leu;
- vi) Arg Tyr Ile Leu Xaa Gly Ile Ser Ala Leu Arg Lys;
- vii) Gly Asp Gln Tyr Xaa Ser Val Leu Met Val Ser Ile;

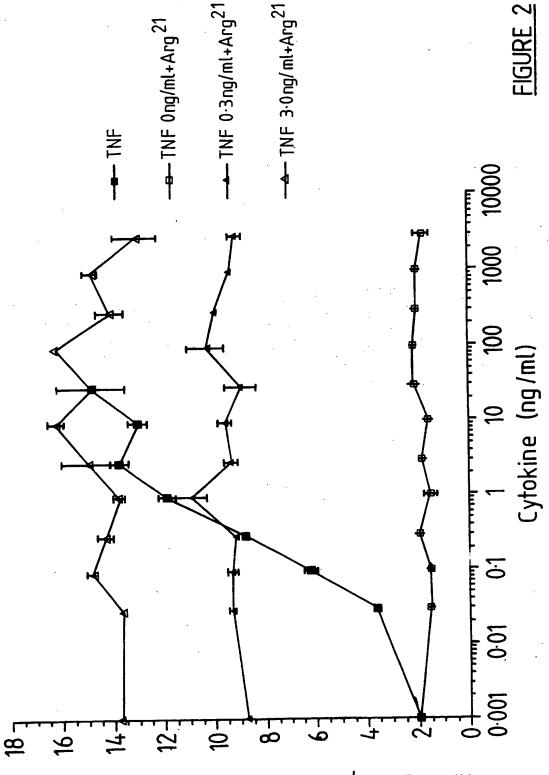
- viii) Ala Gly Ile Leu Xaa Ile Asn Phe Leu Ile Asn Lys Met Gln Glu Asp;
- ix) Asn Met Leu Arg Xaa Leu Arg Asp Ala Phe Ser;
- x) Phe Leu Leu Lys Cys Leu Xaa Gln Val Arg Lys Ile; and
- xi) Tyr Leu Leu Glu Ala Lys Xaa Ala Glu Asn Ile Thr Thr Gly;

wherein Xaa is a basic amino acid selected from the group consisting of Arg and Lys and wherein said variant HGF is in unglycosylated form and acts as an antagonist for at least one property of the corresponding native HGF.

- 34. A method according to claim 32 or 33 wherein the HGF is a modified form of an HGF selected from GM-CSF, IL-2, IL-3, IL-4, IL-5, IL-6, Il-7, IL-9, IL-10, G-CSF, EPO.
- 35. A method according to claim 32 or 33 wherein the modified HGF is a modified human GM-CSF having an amino acid sequence in a first α -helix comprising His Val Asn Ala Ile Gln Arg Ala Arg Arg Leu Leu Asn Leu.
- 36. A method according to claim 32 or 33 wherein the modified HGF is a modified human GM-CSF having an amino acid sequence in a first α-helix comprising His Val Asn Ala Ile Gln Lys Ala Arg Arg Leu Leu Asn Leu.
- 37. Use of one or more modified HGFs each as defined in claim 1 or 7 in the manufacture of a medicament for the treatment of the affects of an aberrant endogenous HGF.
- 38. An agent comprising one or more modified HGFs each as defined in claim 1 or 7 for treating the affects of an aberrant endogenous HGF.
- 39. A pharmaceutical composition comprising one or more modified HGFs each as defined in claim 1 or 7 together with one or more pharmaceutically acceptable carriers and/or diluents.



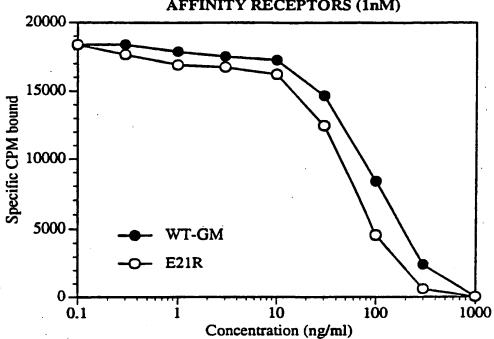
SUBSTITUTE SHEET (Rule 26)



n.moles superoxide \ 1000000 Neutrophils

SUBSTITUTE SHEET (Rule 26)

3/13
COMPETITION OF Wt-GMCSF AND E21R FOR LOW
AFFINITY RECEPTORS (1nM)



COMPETITION OF Wt-GMCSF AND E21R FOR HIGH AFFINITY RECEPTORS (100pM)

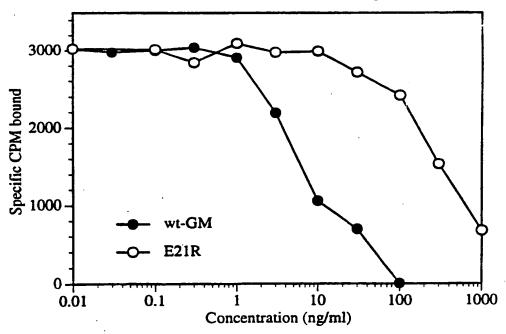
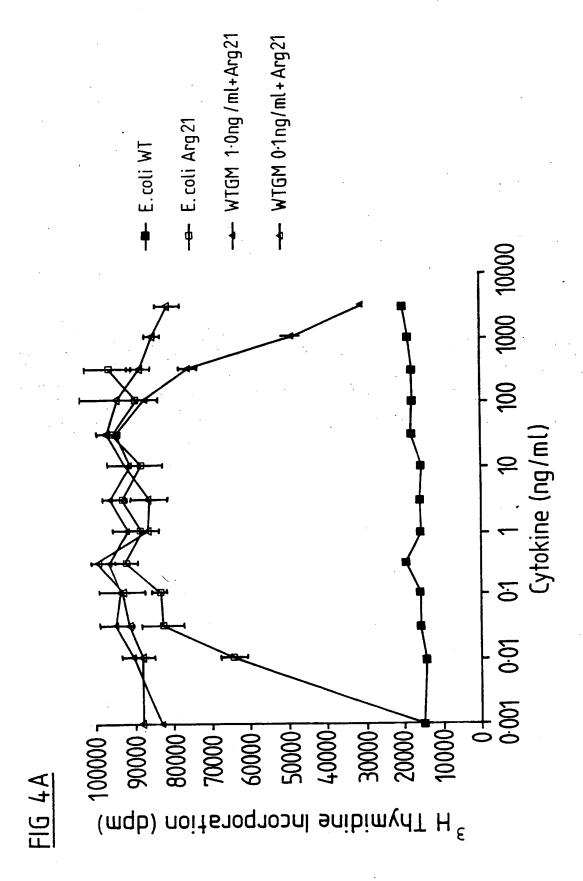
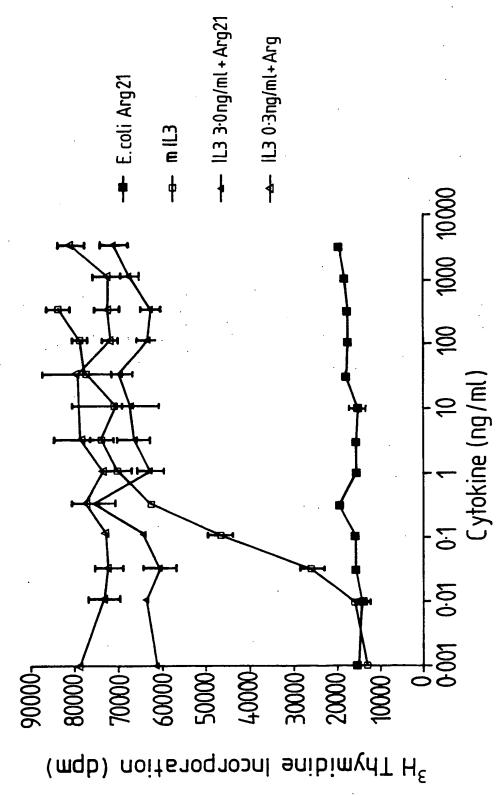


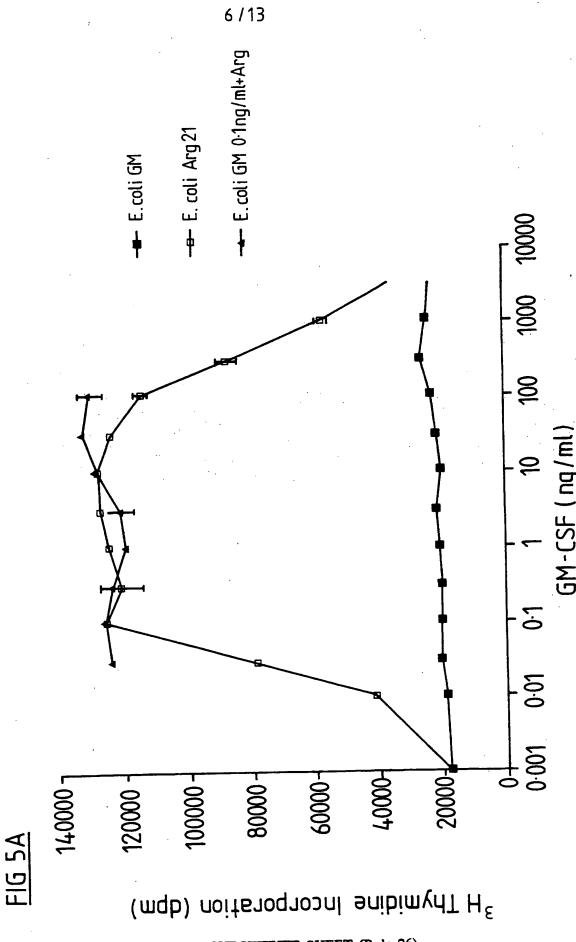
FIGURE 3



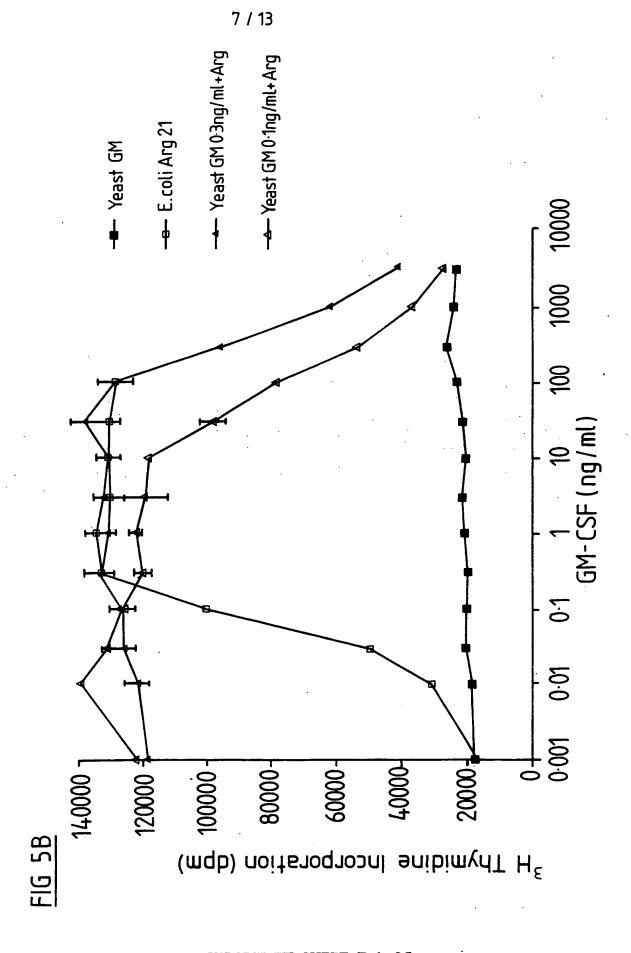
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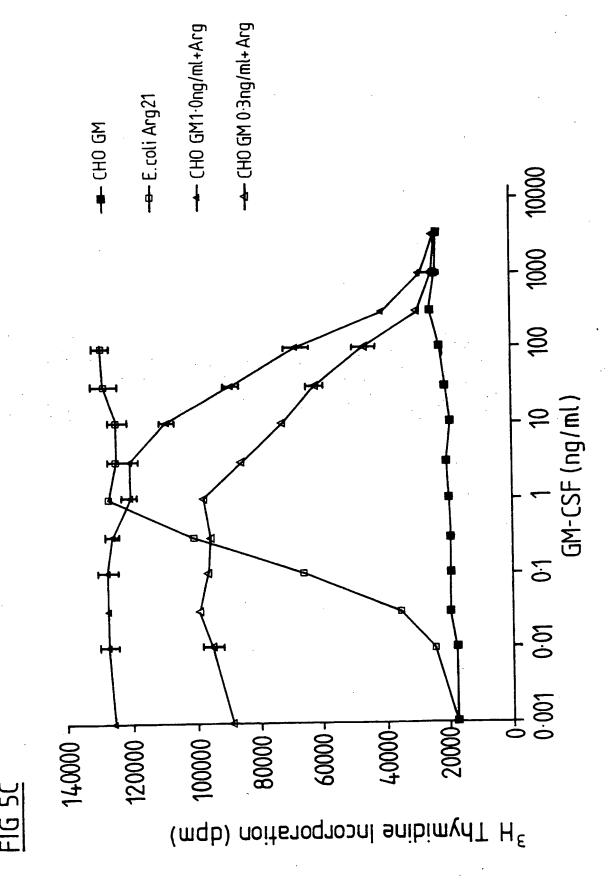


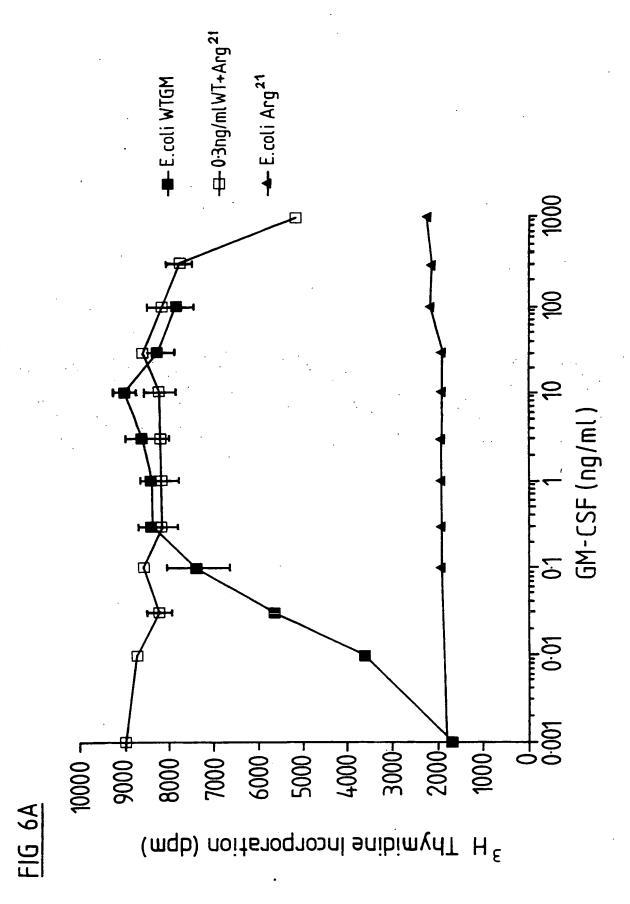
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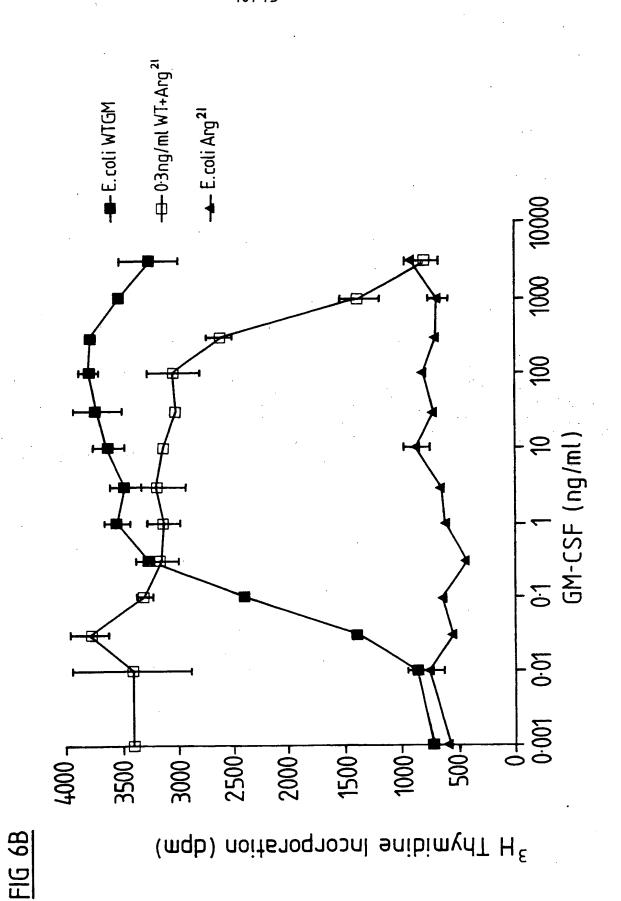
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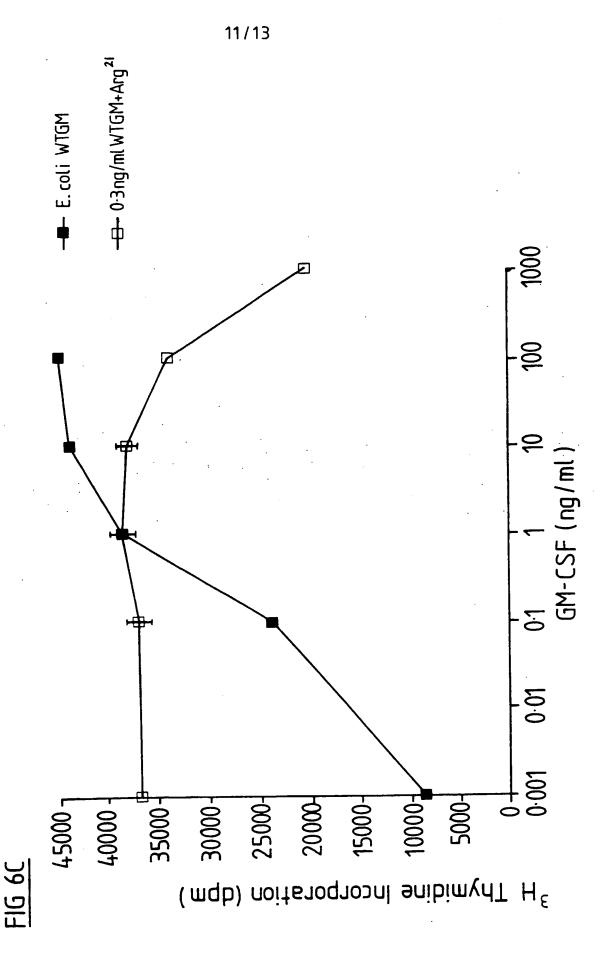




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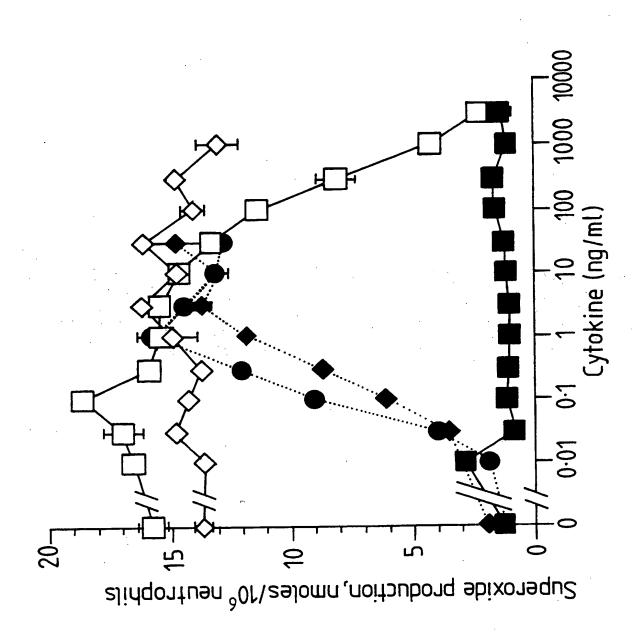
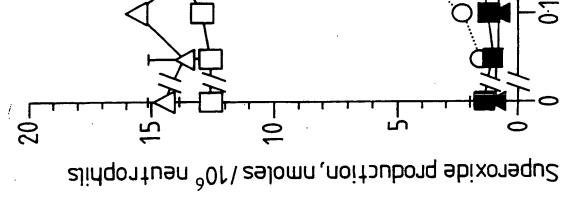
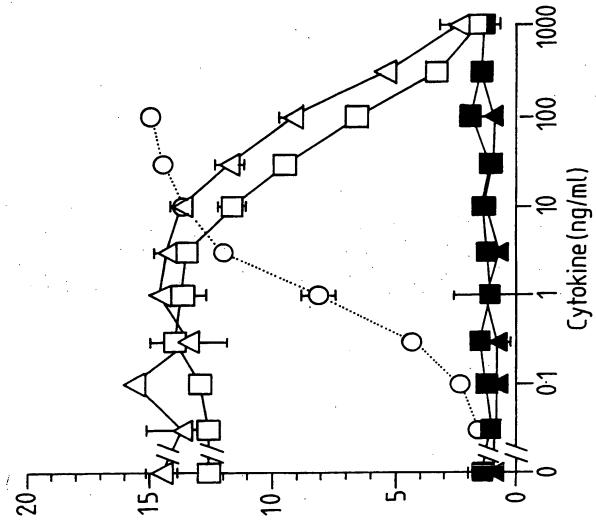


FIG 7A



SUBSTITUTE SHEET (Rule 26)



A. CLASSIFICATION OF SUBJECT MATTER

Int. Cl. 6 C07K 14/52, 14/505, 14/535, 14/54, 14/55, 14/475, C12N 15/19, 15/24, 15/26, 15/27, C12P 21/02, A61K 38/18, 38/19, 38/20

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

Derwent database; World Patent Index (WPAT); American Chemical Society database; Chemical Abstracts (CA). See "electronic data base" box below for key words used.

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched IPC: AU C12N 15/19, 15/24, 15/27, 15/26

Electronic data base consulted during the international search (name of data base, and where practicable, search terms used) WPAT, CA, DERWENT BIOTECHNOLOGY ABSTRACTS DATABASE (BIOT): (KEYWORDS) MUTEIN#; VARIANT#; ANALOG:; COLONY STIMULATING FACTOR#; INTERLEUKIN#; ERYTHROPOIETIN#; HELIC:; HELIX:

Protein sequences - see Supplemental Box

C.	DOCUMENTS CONSIDERED TO BE RELEVA	ANT		· · · · · · · · · · · · · · · · · · ·		
Category*	Citation of document, with indication, where a	Relevant to Claim No.				
x	AU,B,60960/90 (636641) (GIST-BROCADI 14 February 1991 (14.02.91) pages 6, 8, 15, examples 2 and 3	1,2,4-8,15				
x	AU,A,27538/92 (MEDVET SCIENCE PTY page 3, figures 7 and 8	A,27538/92 (MEDVET SCIENCE PTY LTD) 15 April 1993 (15.04.93) 3, figures 7 and 8				
P,X	AU, A, 56709/84 (GD SEARLE & CO.) 9 Ju see pages 191-192, table 6	1,2,4-8,15				
X Furth in the	er documents are listed continuation of Box C.	X	See patent family annex			
-	al categories of cited documents : ment defining the general state of the art which is onsidered to be of particular relevance	υŢu	filing date or priority da with the application but principle of theory und	cited to understand the		
'E" earlie	er document but published on or after the	"X"	invention cannot be cor considered to involve a	relevance; the claimed isidered novel or cannot be n inventive step when the		
	ment which may throw doubts on priority claim(s) nich is cited to establish the publication date of the citation or other special reason (as specified) ment referring to an oral disclosure, use, sition or other means ment published prior to the international filing date the than the priority date claimed	" Y"	inventive step when the	e relevance; the claimed residered to involve an document is combined such documents, such ious to a person skilled in		
		"&"	document member of the	ne same patent family		
Date of the a	ctual completion of the international search		of the international search			
15 Novemb	er 1994 (15.11.94)	17 NO	U 1994 (17.11.94)		
	ailing address of the ISA/AU	Authorized office	er			
AUSTRALIA PO BOX 200 WODEN AO AUSTRALIA	CT 2606	J.H. CHAN				

Telephone No. (06) 2832340

Facsimile No. (06) 2853929

ateg ry	Citation f document, with indication, where appropriate f the relevant passages	Relevant to Claim No.		
P,X	Proceedings of National Academy of Science USA, Vol. 91, issued 21 June 1994, "Specific human granulocyte-macrophage colony-stimulating factor antagonists", (T.R. HERCUS et al), pages 5838-5842, see column 2 page 5839.	1,2,3-12		
P,X	The EMBO Journal, Vol. 12, No. 13 (1993), issued 15 December 1993, "Two distinct functional sites of human interleukin-4 are identified by variants impaired in either receptor binding or receptor activation" (N. KRUSE et al) pages 5121-5129, see table 1 pages 5122-5125.	1,2,4-8,18-31		
P,X	European Journal of Biochemistry, Vol. 222, issued 1 June 1994, "Neutralizing monoclonal antibodies define two different functional sites in human interleukin-4", (REUSCH, P. et al), pages 491-499, see column 1 of page 492, tables 1, 2 and 3.			
x	Proceedings of National Academy of Science USA, Vol. 85, issued October 1988, "Identification of specific residues of human interleukin-2 that effect binding to the 70-KDa subunit (p70) of the interleukin-2 receptor", pages 7709-7713, see tables 1 and 2.	1,2,4-8,16,17		
P,X	Biochemistry 1994, Vol. 33, issued 31 May 1994, "Mutagenic Analysis of a Receptor Contact Site on Interleukin-2: Preparation of an IL-2 Analog with Increased Potency" (BERNDT WILLIAM G. et al), pages 6571-6577, see column 2 page 6571, tables 1 and 2.	1,2,4-8,16,17		
x	The EMBO Journal, Vol. 11, No. 3, issued March 1992, "Residue 21 of human granulocyte-macrophage colony stimulating factor is critical for biological activity and for high but not low affinity binding", (A.F. LOPEZ et al), pages 909-916. See column 1 of page 910 to column 2 page 912.	1,2-12		
A .	Biochemical and Biophysical Research Communications, Vol. 159, No. 1, 1989, issued 28 February 1989, "Mutagenesis of human granulocyte colony stimulating factor", pages 103-111.			
A	Proceedings of National Academy of Science, Vol. 89, issued 1 December 1992, "A human interleukin-3 analog with increased biological and binding activities", (T. KUGO et al), pages 11842-11846.			
A	Biochimica et Biophysica Acta, Vol. 1041 (1990), "Theoretical conformational analysis of a family of alpha-helical immunocytokines (V.P. ZAV'YALOV et al), pages 178-185.			
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PCT/AU 94/00432

B. FIELDS SEARCHED (supplemental box)

The following protein subsequences were searched in CA database in STN International:

HVNAIQ[HKR]ARRLLNL
ALVK[HKR]TLALLSTHRTL
NMI[HKR][DE]IITHL
NMI[DE][HKR]IITHL
NMI[HKR][HKR]IITHL
LLL[HKR]LQMIL
ITLQ[HKR]IIKTL
AYIL[HKR]GISALRK
GDQY[HKR]SVLMVSI
AGIL[HKR]INFLINKMQGD
NMLR[HKR]LADAFS
FLLKCL[HKR]QVRKI

YLLEAK[HKR]AENITTG

END OF ANNEX

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

	Patent Docume Cited in Searc Report				Member					
AU	60690/90		US	5331148	wo	91/03029				
λU	27538/92		EP	609280	wo	93/07171				
\U	56709/94		AU	56125/94	wo	94/12638	wo	94/12639		
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CLAIMS

- A peptide of 5-30 amino acids which peptide exhibits antagonistic activity directed against IL-6.
- A peptide of 5-30 amino acids which peptide exhibits antagonistic activity directed against the α or β -chain of the IL-6 receptor.
- A peptide of 5-30 amino acids which peptide exhibits antagonistic or agenistic IL-6 activity.
- A peptide according to claim 1, 2 or 3 of 5-20 amino
- A peptide according to claim 4 of 5-12 amino acids. 10 5
 - A peptide according to claim 1, 2, 3, 4 or 5 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNI, ILRSFKEPLQSSLRALRQM QLSCFRKSPLSNVVC,
- 150 PRSTPSLTTKAVLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNE-NFTLKSEWATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or pvykvkfnpphnlsvzn
- A peptide according to claim 3, 4 or 5 having at least one string of % consecutive amino acids long in common with the following/amino acid sequence: EWGPRSTPSLTT-20 KAVLLVRKFQNEPAED
 - A pertide composition, wherein at least two pertides according to any of claims 1-7 are chemically linked directly or via spacer molecules.
- A peptide composition according to claim 8 wherein at 25 least two peptides are linked with lysine.
 - A peptide composition wherein at least two peptides according to claim 1, 2, 3, 4, or 5 having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK are linked with lysine.

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A peptide composition according to claim 8, 9 or 10 wherein at least four peptides are linked with branching oligolysines.

A mixture comprising peptides and/or peptide compositions according to any of claims 1-11.

- Antibody specifically directed against a peptide or a peptide composition according to any of claims 1-11.
- Anti-idiot/pic antibody raised against an antibody according to claim 13.
- A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to any of the above claims together with at least one suitable excipient for administration.
- Use of a pharmaceutical preparation according to claim
- 15 in the treatment or prevention of an IL-6 related disease.
 - Use of a peptide, peptide composition or antibody according to anyone of claims 1-13 to clear extra-corporeal blood or blood products from IL-6 or IL-5 receptor molecules.
 - A diagnostic assay comprising a peptide or a peptide composition or an antibody according to anyone of claims
 - Use of a diagnostic assay according to claim 18 to detect or diagnose IL-6 related disease in man or animals.
- Use of a peptide according to claim 7 to exert agonistic IL-6 activity at concentrations that are relatively 25 equivalent to 7.5 to 120 $\mu g/ml$ when tested in vitro in a B9 cell bio assay.
 - Use of a peptide according to claim 20 in cell-culture.
- A pharmaceutical preparation comprising a peptide according to claim 7 together with at least one suitable 30 excipient for administration.
 - Use of a pharmaceutical preparation according to claim 22 for topical or intra-mammary application.
- Use of a peptide, or peptide composition according to anyone of claims 1-12 for the manufacture of a medicament for Multopical of intra-mammary application.

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9/2021/14/CO 63 Rec'd PCT/PTO 18 DEC 1938 PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Loenardus Adrianus Maria Govardus van Leengoed et al.

Serial No.: PCT/NL97/00345

Filed: 19 June 1997

For: IL-6 AND IL-6-RECEPTOR DERIVED PEPTIDES HAVING IL-6 ANTA GONISTIC OR AGONISTIC ACTIVITY

Examiner:

Group Art Unit:

Attorney Docket No.: 3890

NOTICE OF EXPRESS MAILING

Express Mail Mailing Label Number: EL248225768US
Date of Deposit with USPS: 18 December 1998
Person making Deposit:
Jared Turner

PRELIMINARY AMENDMENT

Rox

Assistant Commissioner for Patents Washington, D.C. 20231

Sir:

Please amend the above referenced application as indicated below prior to the calculation of the filing fee.

IN THE CLAIMS:



4. (Amended) A peptide according to claim 1 [, 2, or 3] of 5-20 amino acids.

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6. (Amended) A peptide according to claim 1 [.2,3,4, or 5] having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM, QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNEGKKMRCEWDGGR, NFTLKSEQATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or PVYKVKPNPPHNLSVIN.

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- 7. (Amended) A peptide according to claim 1 [3, 4, or 5] having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLLVRKFQNSPAED.
- 8. (Amended) A peptide composition, wherein at least two peptides according to any of [claims 1-7] claim 1 are chemically linked directly or via spacer molecules.

- 10. (Amended) A peptide composition wherein at least two peptides according to claim 1 [,2, 3, 4, or 5] having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK are linked with lysine.
- 11. (Amended) A peptide composition according to claim 8 [,9 or 10] wherein at least four peptides are linked with branching oligolysines.
- 12. (Amended) A mixture comprising peptides and/or peptide compositions according to [any of claims 1-11] claim 1.
- 13. (Amended) Antibody specifically directed against a peptide or a peptide composition according to [any of claims 1-11] claim 1.

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15. (Amended) A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to [any of the above claims] <u>claim 1</u> together with at least one suitable excipient for administration.

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- 17. (Amended) Use of a peptide, peptide composition or antibody according to [anyone of claims 1-13] <u>claim 1</u> to clear extra-corporeal blood or blood products from IL-6 or IL-6 receptor molecules.
- 18. (Amended) A diagnostic assay comprising a peptide [or a peptide composition or an antibody] according to [anyone of claims 1-13] <u>claim 1</u>.

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24. (Amended) Use of a peptide [, or peptide composition according to anyone of claims 1-12] claim 1 for the manufacture of a medicament for topical or intra-mammary application.

Please add the following new claims:

- 25. A peptide according to claim 2 of 5-20 amino acids.
- 26. A peptide according to claim 25 of 5-12 amino acids.

27. A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence:

EWGPRSTPSLTTKAVLLLVRKFQNSPAED.

- 28. A peptide according to claim 2 of 5-20 amino acids.
- 29. A peptide according to claim 28 of 5-12 amino acids.
- 30. A peptide according to claim 2 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM, QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNLSCIVNEGKKMRCEWDGGR, NFTLKSEQATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or PVYKVKPNPPHNLSVIN.

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31. A peptide according to claim 2 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence:

EWGPRSTPSLTTKAVLLLVRKFQNSPAED.

- 32. A peptide composition, wherein at least two peptides according to claims 2 are chemically linked directly or via spacer molecules.
- 33. A peptide composition according to claim 32 wherein at least two peptides are linked with lysine.
- 34. A peptide composition according to claim 32 wherein at least four peptides are linked with branching oligolysines.
- 35. A peptide composition wherein at least two peptides according to claim 2 having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK are linked with lysipe.
- 36. A mixture comprising peptides according to claim 2.
- 37. Antibody specifically directed against a peptide according to claim 2.
- 38. Anti-idiotypic antibody raised against an antibody according to claim 37.
- 39. A pharmaceutical preparation comprising a peptide according to claim 2 together with at least one suitable excipient for administration.
- 40. Use of a pharmaceutical preparation according to claim 39 in the treatment or prevention of an IL-6 related disease.
- 41. Use of a peptide according to claim 2 to clear extra-corporeal blood or blood products

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from IL-6 or IL-6 receptor molecules.

- 42. A diagnostic assay comprising a peptide according to claim 2.
- 43. Use of a diagnostic assay according to claim 42 to detect or diagnose IL-6 related disease in man or animals.
- 44. Use of a peptide according to claim 31 to exert agonistic IL-6 activity at concentrations that are relatively equivalent to 7.5 to 120 micrograms/ml when tested in vitro in a B9 cell bio assay.
- 45. Use of a peptide according to claim 44 in a cell-culture.
- 46. A pharmaceutical preparation comprising a peptide according to claim 31 together with at least one suitable excipient for administration.
- 47. Use of a pharmaceutical preparation according to claim 6 for topical or intra-mammary application.
- 48. Use of a peptide according to claim 2 for the manufacture of a medicament for topical or intra-mammary application.
- 49. A peptide according to claim 3 of 5-20 amino acids.
- 50. A peptide according to claim 49 of 5-12 amino acids.
- 51. A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL, ILRSFKEFLQSSLRALRQM, QLSCFRKSPLSNVVC, PRSTPSLTTKAVLLLVRKFQNS, MCVASSVGSKFSKTQTFQGC, PEKPKNVSCIVNEGKKMRCEWDGGR,

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NFTLKSEQATHKFADCKAKRDTPTS, WVEAENALGKVTSDH, or PVYKVKPNPPHNLSVIN.

- 52. A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence:

 EWGPRSTPSLTTKAVLLLVRKFQNSPAED.
- 53. A peptide composition, wherein at least two peptides according to claim 3 are chemically linked directly or via spacer molecules.
- 54. A peptide composition according to claim 53 wherein at least two peptides are linked with lysine.
- 55. A peptide composition wherein at least two peptides according to claim 3 having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK are linked with lysine.
- 56. A peptide composition according to claim 53 wherein at least four peptides are linked with branching oligolysines.
- 57. A peptide composition according to claim 55 wherein at least four peptides are linked with branching oligolysines.
- 58. A mixture comprising peptides and/or peptide compositions according to claim 3.
- 59. Antibody specifically directed against a peptide or a peptide composition according to claim 3.
- 60. Anti-idiotypic antibody raised against an antibody according to claim 59.

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- 61. A pharmaceutical preparation comprising a peptide or a peptide composition or an antibody according to claim 3 together with at least one suitable excipient for administration.
- 62. Use of a pharmaceutical preparation according to claim 61 in the treatment or prevention of an IL-6 related disease.
- 63. Use of a peptide, peptide composition or antibody according to claim 3 to clear extracorporeal blood or blood products from IL-6 or IL-6 receptor molecules.
- 64. A diagnostic assay comprising a peptide according to claim 3.
- 65. Use of a diagnostic assay according to claim 64 to detect or diagnose IL-6 related disease in man or animals.
- 65. Use of a peptide according to claim 3 for the manufacture of a medicament for topical or intra-mammary application.
- 66. Use of a peptide according to claim 52 to exert agonistic IL-6 activity at concentrations that are relatively equivalent to 7.5 to 120 micrograms/ml when tested in vitro in a B9 cell bio assay.
- 67. Use of a peptide according to claim 66 in a cell-culture.
- 68. A pharmaceutical preparation comprising a peptide according to claim 52 together with at least one suitable excipient for administration.
- 69. Use of a pharmaceutical preparation according to claim 68 for topical or intra-mammary application.
- 70. Use of a peptide according to claim 3 for the manufacture of a medicament for topical

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REMARKS

The Office is respectfully requested to enter the above amendment prior to the calculation of the filing fee.

Respectfully submitted,

Laurence B. Bond

Registration No. 30,549

Attorney for Applicants

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Salt Lake City, Utah 84110-2550

Telephone: (801) 532-1922

Date: December 18, 1998

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On page 15, line 27, after "PAED" please insert --(SEQ. ID. NO.: 11)--;
On page 16, line 5, after "SVIN" please insert --(SEQ. ID. NO.: 10)--;
On page 16, line 5, after "TSDH" please insert --(SEQ. ID. NO.: 9)--;
On page 16, line 5, after "FQGC" please insert --(SEQ. ID. NO.: 6)-.

IN THE FIGURES

NR.

On FIG. 1, after "SSLRALRQM" please insert --(SEQ. ID. NO.: 12)--;

On FIG. 2, after "PPANITV" please insert -- (SEQ. ID. NO.: 13)--;

On FIG. 3, after "PHNLSVIN" please insert --(SEQ. ID. NO.: 14)-.

IN THE CLAIMS:

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- 6. (Twice Amended) A peptide according to claim 1 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL (SEQ. ID. NO.: 2), ILRSFKEFLQSSLRALRQM (SEQ. ID. NO.: 3), QLSCFRKSPLSNVVC (SEQ. ID. NO.: 4), PRSTPSLTTKAVLLLVRKFQNS (SEQ. ID. NO.: 5), MCVASSVGSKFSKTQTFQGC (SEQ. ID. NO.: 6), PEKPKNLSCIVNEGKKMRCEWDGGR (SEQ. ID. NO.: 7), NFTLKSEQATHKFADCKAKRDTPTS (SEQ. ID. NO.: 8), WVEAENALGKVTSDH (SEQ. ID. NO.: 9), or PVYKVKPNPPHNLSVIN (SEQ. ID. NO.: 10).
 - 7. (Twice Amended) A peptide according to claim 1 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLLVRKFQNSPAED (SEQ. ID. NO.: 11).



10. (Twice Amended) A peptide composition wherein at least two peptides according to claim 1 having at least one string of 5 consecutive amino acids long in common with the amino acid

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sequence RYILDGISALRK (SEQ. ID. NO.: 1) are linked with lysine.

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27. (Amended) A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLLVRKFQNSPAED (SEQ. ID. NO.: 11).

BY

30. (Amended) A peptide according to claim 2 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL (SEQ. ID. NO.: 2), ILRSFKEFLQSSLRALRQM (SEQ. ID. NO.: 3), QLSCFRKSPLSNVVC (SEQ. ID. NO.: 4), PRSTPSLTTKAVLLLVRKFQNS (SEQ. ID. NO.: 5), MCVASSVGSKFSKTQTFQGC (SEQ. ID. NO.: 6), PEKPKNLSCIVNEGKKMRCEWDGGR (SEQ. ID. NO.: 7), NFTLKSEQATHKFADCKAKRDTPTS (SEQ. ID. NO.: 8), WVEAENALGKVTSDH (SEQ. ID. NO.: 9), or PVYKVKPNPPHNLSVIN (SEQ. ID. NO.: 10).

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31. (Amended) A peptide according to claim 2 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLLVRKFQNSPAED (SEQ. ID. NO.: 11).

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35. (Amended) A peptide composition wherein at least two peptides according to claim 2 having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK (SEO. ID. NO.: 1) are linked with lysine.

B7

51. (Amended) A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with one of the following amino acid sequences: STKVLIQFLQKKAKNL (SEQ. ID. NO.: 2), ILRSFKEFLQSSLRALRQM (SEQ. ID. NO.: 3), QLSCFRKSPLSNVVC (SEQ. ID. NO.: 4), PRSTPSLTTKAVLLLVRKFQNS (SEQ. ID. NO.: 5), MCVASSVGSKFSKTQTFQGC (SEQ. ID. NO.: 6), PEKPKNLSCIVNEGKKMRCEWDGGR (SEQ. ID. NO.: 7), NFTLKSEQATHKFADCKAKRDTPTS (SEQ. ID. NO.: 8), WVEAENALGKVTSDH (SEQ. ID. NO.: 9), or PVYKVKPNPPHNLSVIN (SEQ. ID. NO.: 10).

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Conclude

52. (Amended) A peptide according to claim 3 having at least one string of 5 consecutive amino acids long in common with the following amino acid sequence: EWGPRSTPSLTTKAVLLLVRKFQNSPAED (SEQ. ID. NO.: 11).

BY

55. (Amended) A peptide composition wherein at least two peptides according to claim 3 having at least one string of 5 consecutive amino acids long in common with the amino acid sequence RYILDGISALRK (SEQ. ID. NO.: 1) are linked with lysine.

REMARKS

The application is to be amended without prejudice or disclaimer as previously set forth.

Primarily, the amendments are sought to conform the application to a form more consistent with

Office practice.

The application is also to be amended to include a substitute SEQUENCE LISTING in conformity with 37 C.F.R. §§ 1.821-1.825. Submitted herewith is the substitute SEQUENCE LISTING, a copy of the substitute SEQUENCE LISTING in computer readable form, as well as a Statement Under 37 C.F.R. §§ 1.821(f) and 1.825.

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